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A00432 Nucleotide (A25812 polynucleot A40518 Sequence 55 A88271 Sequence 41 A88271 Sequence 42 A890238 Sequence 41 A90238 Sequence 41 A90240 Sequence 42 AR119028 Sequence AX007733 Sequence AX007734 Sequence AX007734 Sequence

AR062865 Sequence AX073580 Sequence

Description

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Microfabricated, flowthrough porous apparatus detection of binding reactions Patent: US 5843767-A 13 01-DEC-1998;
             Beattie, K.L.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

BASE COUNT ORIGIN

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/organism="unknown" 4 c 0 g Location/Qualifiers

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5843767.

PAT

29-SEP-1999

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Matches 7; Conservative 0; Mismatches 0;
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1 (bases 1 to 12)

Schmidt,A.C., Składopoulos,M.H., Collins,P.L., Murphy,B.R.,
Bailly,J.E. and Durbin,A.P.
Attenuated human-bovine chimeric parainfluenza virus (piv)
Patent: WO 0104330-A 2 18-JAN-2001;
THE GOVERNMENT OF THE UNITED STATES OF AMERICA (US)
LOCation/Qualifiers
                                                                                                                                                                                                  unidentified.
unidentified
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                                                                                                                              PHARMACEUTICAL COMPOSITION USEFUL IN THE PAPILLOMAVIRUS-INDUCED TUMOURS Patent: WO 9010459-A 7 20-SEP-1990; Transgene S.A
                                                                                                                                                                                                                                                          A00432 13 bp
Nucleotide sequence A00432
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/note="Sequence flanking site for introduction of Sgr Al
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Brysch, W. and Schlingensiepen, K.
AN ANTISENSE OLIGONUCLEOTIDE PREPARATION METHOD
                                                                                                                                                                                                                                                                                                                                  1 (bases 1 to 14)
Brysch,W. and Schlingensiepen,K.
AN ANTISENSE OLIGONUCLEOTIDE PREPARATION METHOD
PATENT: WO 9833904-A 421 06-AUG-1998;
BIOGNOSTIK GES (DE); BRYSCH WOLFGANG (DE)
Location/Qualifiers
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Brysch, W. and Schlingensiepen, K.
ANTISENSE OLIGONUCLEOTIDE PREPARATION METHOD Patent: WO 9833904-A 419 06-AUG-1998;
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Erysch.W.D. and Schlingensiepen.K.D.
An antisense oligonucleotide preparation method
Patent: ED 0856579-A 419 05-AUG-1998;
BIOGNOSTIK GES (DE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Patent: WO 9833904-A 1193 06-AUG-1998;
BIOGNOSTIK GES (DE); BRYSCH WOLFGANG (
Location/Qualifiers
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Sequence 421 from Patent
A90240 Transfer
                                                                     1 (bases 1 to 14)
Brysch,W.D. and Schlingensiepen,K.D.
An antisense oligonucleotide preparation method
Patent: EP 0856579-A 421 05-AUG-1998;
                                                                                                                     unidentified unclassified
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                                                                BIOGNOSTIK GES (DE)
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      /organism="unidentified"
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AR119028
AR119028.1
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Sequence 275 from Patent W09967428.
AX007733
AX007733.1 GI:9995430
                                                                                                                                                              Patent: WO 9967428-A 275 29-DEC-1999; INNOGENETICS NV (BE); STUYVER LIEVEN
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Uchida, X., Uchida, T., Tanaka, Y., Matsuda, Y. and Kondo, S.
Antisense nucleic acid compound targeted to VEGF
Patent: US 6150092-A 154 21 NOV-2000;
                                                                                                                                                                                           Method for detection of drug-selected mutations in the hiv protease
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Viruses; Retroid viruses;
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154 from patent
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Pred. No. 5e+
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1 (bases 1 to 14)

Brysch,W., Schlingensiepen,K.H. and Schlingensiepen,R.

A method for stimulating the immune system Patent: WO 963975-A 43 16-DEC-1999;

BIOGNOSTIK GES (DE); BRYSCH WOLFGANG (DE); SCHLINGENSIEPEN KARL HERMANN (DE); SCHLINGENSIEPEN REIMAR (DE)
Sequence 19 from patent 126234 1 GI:1606104
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AX007734
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INNOGENETICS NV (BE); STUYVER LIEVEN (BE)
Location/Qualifiers
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/db_xref="taxon:11966"
6 c 0 g 3 t
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KEYWORDS
SOURCE
ORGANISM Unknown.

REFERENCE 1 (bases 1 to 14)
AUTHORS
Vergnaud,G.
TITLE Process for detection of new polymorphic loci in a DNA sequence, nucleotide sequences forming hybridization probes and their
JOURNAL Patent: US 556955-A 19 17-SEP-1996;
FEATURES 1.14

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Ouery Match 100.0%; Score 7; DB 6; Length 14;
Best Local Similarity 100.0%; Pred. No. 5e+05;
Matches 7; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Ouery Total Similarity 100.0%; Pred. No. 5e+05;
Matches 7; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Metastatic breast
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                                                                                                                                                      Description
                                                                                                     Leptospira nucleot
Human parkin gene
Human dendritic ce
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ALIGNMENTS

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New isolated pathogenic Leptospira bacterium - useful for, developing products for conferring protective immunity, and prophylactic or therapeutic treatment
                                                                              WPI; 1998-520791/44.
                                                                                                                                                        (AGRI-) AGRIC VICTORIA SERVICES (PIGR-) PIG RES & DEV CORP.
                                                                                                                                                                                                                    07-MAR-1997;
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                                                                                                                                                                                                                                                                                                                                                                             Leptospira sp.
                                                                                                                                                                                                                                                                                                                                                                                                                     Infection; pathogenic Leptospira; protective immunity; therapy; diagnosis; ss.
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Claim 15; Page 71; 94pp; English.

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Best Local
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                                                                                                                 This sequence represents a fragment of an intron from the gene of the invention. The gene has been designated the parkin gene, and variants it are implicated in the pathology of Parkinson's disease, and found i parkinson's disease patients. The sequences may be used for the diagnosis, treatment (including gene therapy) and investigation of
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Parkinson's disease related gene; parkin gene; variant; gene therapy; intron; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   diagnosis of past or present LS infection.
                                                                                                                                                                                         Claim 11; Page 100; 114pp; English.
                                                                                                                                                                                                                Gene implicated in the pathology of Parkinson's disease, treatment of the disease
                                                                                                                                                                                                                                                   WPI; 1999-494295/41.
                                                                                                                                                                                                                                                                          Mizuno Y,
                                                                                                                                                                                                                                                                                                                                     09-FEB-1998;
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                                                                                Sequence 10
                                                                                                                                                                                                                                                                                                (MIZU/) MIZUNO Y. (SHIM/) SHIMIZU N
                                                                                                                                                                                                                                                                                                                                                                                   12-AUG-1999
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                                                                                                          Parkinson's disease.
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RESULT
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19-JUN-1998;
19-JUN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human dendritic cell SAGE tag,
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Sequences AA277573-Z79709 represent SAGE (serial analysis of gene expression) tags used to identify mRNA transcripts encoding immunostimulatory cofactor proteins which are preferentially or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cytotoxic
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                                        Claim 1;
                                                        Isolated polynucleotides differentially expressed in cells, useful in gene vaccines against cancer \dot{\ }
                                                                                                                                                                   08-DEC-1998;
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19-JUN-1998;
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| TGTTGGA 3
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                                                                                                                                     GENZYME CORP.
                                                                                                                             SHANKARA
                                        Page 84; 130pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          T-lymphocyte; tumour antigen; immunotherapy; anticancer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                          Shankara
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98US-0090041.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
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                                                                                                                                                           19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
(GENZ ) GENZYME CORP
(ROBE/) ROBERTS B L.
(SHAN/) SHANKARA S.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 antimetastatic;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; metastatic breast tumour tissue; breast cancer; tag; primer;
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98US-0090040.
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vaccine; diagnosis; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Α,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4 T; 0 other;
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hes 0;
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RESULT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; metastatic breast tumour tissue; breast cancer; tag; primer; non-metastatic breast tumour tissue; gene therapy; anticancer; antimetastatic; vaccine; diagnosis; ss.
                                                         19-JUN-1998;
19-JUN-1998;
                                                                                                                              19-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Metastatic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        07-APR-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAZ82660 standard; DNA; 10 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               breast tumour tissue (i.e. are downregulated in metastatic breast tumour
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1; Page 63; 219pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Isolated polynucleotides differentially expressed between metastatic
and non-metastatic breast cancer cells, useful for diagnosis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Roberts BL,
                                                                                                                                                                                                                             18-JUN-1999;
                                                                                                                                                                                                                                                                                              23-DEC-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      immunotherapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           breast
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Shankara
                                                         98US-0089853.
98US-0089997.
98US-0090039.
98US-0090040.
                                                                                                                                                                                                                             99WO-US13647.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       tumour cell upregulated transcript tag #1894.
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Pred. No. 3.3e+04;
; Mismatches 0;
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RESULT
AAZ84297
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Best Local Similarity
Watches 7; Conserv
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19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               promoters from the transcripts are used to direct expression, in selected cell types, of e.g. therapeutic genes (also ribozymes or antisense sequences), particularly an antigen-encoding sequence for use in gene or cell-based vaccines. Polypeptides encoded by the transcripts are also useful in vaccines; for diagnosing breast cancer and for raising specific antibodies (Ab). Ab are used to detect the polypeptides or as therapeutic agents. Host cells that produce the polypeptides can be used to expand and isolate populations of educated, antigen-specific immune effecter cells, e.g. cytotoxic T lymphocytes, and these used for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAZ80767 to AAZ83941 represent tags corresponding to distinct transcripts that are preferentially transcribed in the metastatic breast tumour tissue (i.e. are upregulated in metastatic breast tumour cells).

AAZ83942 to AAZ86677 represent tags corresponding to distinct transcripts that are preferentially transcribed in the primary or non-metastatic breast tumour tissue (i.e. are downregulated in metastatic breast tumour cells). These transcripts can be used for diagnosis, prognosis, monitoring and treatment of breast cancer, particularly where metastatic. Diagnosis is by standard immunoassays or hybridisation/amplification reactions. Compounds that modulate expression of the transcripts are potentially useful for treatment of (metastatic) breast cancer, while
                                                                                                                                                                                              antimetastatic;
                                                                                                                                                                                                                                                            Metastatic breast tumour cell downregulated transcript tag #3531.
                                                                                                                                                                                                                                                                                              07-APR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; Page 110; 219pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Isolated polynucleotides differentially and non-metastatic breast cancer cells, prevention and treatment of cancer -
                                                                                                23-DEC-1999
                                                                                                                                WO9965928-A2
                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                               non-metastatic
                                                                                                                                                                                                                            Human; metastatic breast tumour tissue; breast cancer; tag; primer;
                                                                                                                                                                                                                                                                                                                              AAZ84297;
                                                                                                                                                                                                                                                                                                                                                             AAZ84297 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 10
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                                                                18-JUN-1999;
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2 tgttgga 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   immunotherapy.
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98US-0089853.
98US-0089997.
98US-0090039.
                                                                                                                                                                                                            breast tumour tissue; gene therapy; anticancer;
                                                                99WO-US13647
                                                                                                                                                                                              vaccine; diagnosis;
                                                                                                                                                                                                                                                                                                                                                             DNA;
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 10;
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AAZ80767 to AAZ83941 represent tags corresponding to distinct transcripts that are preferentially transcribed in the metastatic breast tumour tissue (i.e. are upregulated in metastatic breast tumour cells). AAZ83942 to AAZ86677 represent tags corresponding to distinct transcripts that are preferentially transcribed in the primary or non-metastatic breast tumour tissue (i.e. are downregulated in metastatic breast tumour cells). These transcripts can be used for diagnosis, prognosis, monitoring and treatment of breast cancer, particularly where metastatic. Diagnosis is by standard immunoassays or hybridisation/amplification reactions. Compounds that modulate expression of the transcripts are
                                                          potentially useful for treatment of (metastatic) breast cancer, while promoters from the transcripts are used to direct expression, in select cell types, of e.g. therapeutic genes (also ribozymes or antisense sequences), particularly an antigen-encoding sequence for use in gene cell-based vaccines. Polypeptides encoded by the transcripts are also useful in vaccines; for diagnosing breast cancer and for raising specific antibodies (Ab). Ab are used to detect the polypeptides or as therapeutic agents. Host cells that produce the polypeptides can be us to expand and isolate populations of educated, antigen-specific immune effecter cells, e.g. cytotoxic T lymphocytes, and these used for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1; Page 153; 219pp; English.
Sequence 10
                                        adoptive immunotherapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 prevention and treatment of cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Isolated polynucleotides differentially expressed between metastatic and non-metastatic breast cancer cells, useful for diagnosis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2000-106079/09
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(ROBE/)
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19-JUN-1998;
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ROBERTS B L.
BP; 2 A; 0 C;
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98US-0090041.
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  3 G; 5 T; 0 other;
                                                                               e polypeptides can be use antigen-specific immune
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RESULT
AAH18992
ID AAH1
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XX AAH1
XX UCP3
XX UCP
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Best Local S
Matches 7
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                                                                                08-SEP-2000; 2000WO-US24784.
                                                                                                                                                                                                                                                                                                                                                                                                                UCP3; uncoupling protein 3; polymorphism; obesity;
diabetes mellitus; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        21-JUN-2001
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nes 7; Conser
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99US-0152789
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 21; 1
3.3e+04;
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RESULT 8
AAF35297/c
ID AAF35297 standard; DNA; 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention relates to the human uncoupling protein 3 (mitochondrial, proton carrier) (UCP3) gene and polymorphisms. The polymorphisms are associated with obesity, especially diabetes mellitus associated obesity. They polymorphisms may be identified and analysed to determine whether an individual is susceptible to obesity and may be used as the basis for targeted design of drugs to treat obesity. The present sequence was used the identification and amplification of UCP3 polymorphisms.
                                                                                                                                                                                                                                                                                                                nor previously assigned open reading frame; nonannotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification; linker; PCR primer; ds.
                                                                                                                 Velculescu V, Vogelstein·B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nucleic acids encoding uncoupling protein 3 (mitochondrial, proton carrier) (UCP3) proteins comprising single nucleotide polymorphisms useful for the design of drugs for treating obesity -
                                                               Yeast gene coding sequences comprising of gene expression (SAGE) tags, useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Chew A,
                       Example; Page 72; 419pp; English.
                                                 affecting phases of the cell cycle
                                                                                                     WPI; 2001-061874/07.
                                                                                                                                                                                   16-JUN-1999;
                                                                                                                                                                                                           14-JUN-2000; 2000WO-US16223
                                                                                                                                                                                                                                                               WO200077214-A2
                                                                                                                                                                                                                                                                                         Saccharomyces cerevisiae
                                                                                                                                                                                                                                                                                                                                                        Yeast; Saccharomyces cerevisiae; characterisation;
                                                                                                                                                                                                                                                                                                                                                                                  Yeast NORF gene SAGE tag oligonucleotide SEQ ID NO: 2036.
                                                                                                                                                                                                                                                                                                                                                                                                           23-MAR-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                        AAF35297;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Page 23; 94pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2001-218562/22
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                                                                                                                                                        (UYJO ) UNIV JOHNS HOPKINS.
                                                                                                                                                                                                                                      21-DEC-2000.
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1 tgttgga 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Choi JY, Denton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%; (ilarity 100.0%; (Conservative 0;
                                                                                                                                                                                  9905-0335032
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Pred. No. 3.3e+04;
; Mismatches 0;
                                                                                                                               Kinzler
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                                                             NORF genes with serial analysis for studying, monitoring and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 10,
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                                                                                                                                                                                                                                                                                                                                                        cell cycle; NORF;
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The present invention describes an isolated DNA molecule comprising

Yeast gene coding sequences comprising NORF genes with serial analysis of gene expression (SAGE) tags, useful for studying, monitoring and

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RESULT 9
AAF42210/c
ID AAF42210 standard; DNA; 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          expression varies as in M1, where a test substance which modifies the corporession of the yeast gene is a candidate antifungal drug; (3) a method (M3) for identifying human genes which are involved in cell cycle progression comprising contacting human DNA with a probe which comprises at least 10 contiguous nucleotides of a NORF gene whose expression varies as in M1; and (4) a method (M4) for identifying a candidate drug as a member of a class of drugs having a characteristic effect on gene expression in a yeast cell comprising contacting a yeast cell with a candidate drug and monitoring expression in the yeast cell of at least 1 candidate drug and monitoring expression in the yeast cell of at least 1 condidate drug and monitoring expression in the yeast cell of each case of the cell cycle. The methods may be used as markers of phases of the cell cycle. The methods may be used to identify candidate drugs which affect the cell cycle and for identification of antifungal drugs.

AAF33268 to AAF44064 represent SAGE tags used in the exemplification of the present invention. AAF33267 to AAF33267 to represent linkers and PCR primers used in the SAGE method, in the exemplification of the present
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Best Local
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                                                                                                                                                                                                                                                                                                                                      Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF; nor previously assigned open reading frame; nonannotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification; linker; PCR primer; ds.
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                                                    WPI; 2001-061874/07
                                                                                     Velculescu V,
                                                                                                                                                                                              14-JUN-2000; 2000WO-US16223
                                                                                                                                                                                                                                                                   WO200077214-A2
                                                                                                                                                                                                                                                                                                       Saccharomyces cerevisiae
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                                                                                                                     (UYJO ) UNIV JOHNS HOPKINS
                                                                                                                                                            16-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                Yeast NORF gene SAGE tag oligonucleotide SEQ ID NO:8949
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es 7; Conserv
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                                                                                    Vogelstein B,
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Pred. No.
                                                                                     Kinzler
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3.3e+04;
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affecting phases of the cell cycle -

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CC AAF33268 to AAF44064 represent SAGE tags used in the exemplification of the present invention. AAF33262 to AAF33267 represent linkers and PCR convention.
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Best Local :
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                                                                                                                                                                                                                                                                   nor previously assigned open reading frame; nonannotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification; linker; PCR primer; ds.
                                                                                                          14-JUN-2000; 2000WO-US16223
                                                                                                                                                                                                                                Saccharomyces cerevisiae
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Velculescu V,
                                      (UYJO ) UNIV JOHNS HOPKINS.
                                                                                                                                                                                              WO200077214-A2
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| 10 TGTTGGA 4
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nes 7; Conserv
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ilarity 100.0%;
Conservative (
Vogelstein B,
                                                                           99US-0335032
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Pred. No.
Kinzler K;
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RESULT 11 AAX77658/c ID AAX77658 standard; DNA; 11 BP

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30-MAR-1998; 21-NOV-1997; 20-NOV-1998; 03-JUN-1999 W09927135-A2 Synthetic

98WO-US24854.

External guide sequence; EGS;
inactivation; essential gene;

target mRNA;
therapy; ss.

identification; diagnostic;

N11 active

EGS

22.

09-AUG-1999 (first entry)

AAX77658;

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Query Match
Best Local Similarity
Thes 7; Conserva
     g
                                              Ωy
                                                                                                                                                                                                                               CC antifungal drugs comprising: (a) contacting a test substance with a CC yeast cell; and (b) monitoring expression of a NORF gene whose CC expression varies as in M1, where a test substance which modifies the CC expression of the yeast gene is a candidate antifungal drug; (3) a method CC (M3) for identifying human genes which are involved in cell cycle CC progression comprising contacting human DNA with a probe which comprises CC as tast loo contiguous nucleotides of a NORF gene whose expression varies CC as in M1; and (4) a method (M4) for identifying a candidate drug as a CC member of a class of drugs having a characteristic effect on gene CC expression in a yeast cell comprising contacting a yeast cell of at least 1 CC candidate drug and monitoring expression in the yeast cell of at least 1 CC candidate drug and monitoring expression in the yeast cell of at least 1 CC configure whose expression is affected by the class of drugs. The NORF CC genes may be used to study, monitor and affect phases of the cell cycle, the differentially expressed genes may be used as markers of phases of the cell cycle. The methods may be used to identify candidate drugs which caffect the cell cycle and for identification of antifungal drugs.

CC AAF33268 to AAF44064 represent SAGE tags used in the exemplification of the present invention. AAF33262 to AAF33267 represent linkers and PCR comprisers used in the SAGE method, in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention describes an isolated DNA molecule comprising a coding sequence of a yeast gene selected from a group of 745 NORF (not previously assigned open reading frame, or nonannotated ORF) genes comprising a SAGE (serial analysis of gene expression) tag. Also described are: (1) a method (M1) of using NORF genes to affect the cell cycle comprising administering a NORF gene whose expression varies by at least 10% between any two phases of the cell cycle selected from log phase. S phase and G2/M; (2) a method (M2) for screening candidate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Yeast gene coding sequences comprising of gene expression (SAGE) tags, useful affecting phases of the cell cycle -
                                                                                                                                                                                                                          Sequence 10 BP; 1 A; 0 C; 6
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1 tgttgga 7
                       1 tgttgga 7
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                                                                                                 Conservative
                                                                                                                        100.0%;
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Pred. No.
                                                                                               Mismatches
                                                                                                                        DB 22;
. 3.3e+04;
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Parainfluenza virus; PIV; infectious; human; nucleocapsid protein; BPIV; nucleocapsid phosphoprotein; polymerase protein; human; bovine; HPIV; chimeric; vaccine; immune response; HPIV1; HPIV2; HPIV3; BPIV3; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BPIV3 nucleotide fragment.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cells
                                                                                                          Schmidt AC,
Durbin AP;
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                                        WPI; 2001-081053/09
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                                                                                                                                                                                                             DEPT HEALTH & HUMAN SERVICES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        B₽;
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                                                                                                                                           Skiadopoulos MH,
                                                                                                                                                                                                                                                                                  99US-0143134.
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100.0%; Pr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             58pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Robertson HD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 7;
Pred. No.
                                                                                                                                           Collins PL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 20; 1
                                                                                                                                       Murphy BR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0,
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                                                                                                                                           Bailly JE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0;
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RESULT 13
AAQ40606/c
ID AAQ406
XX AAQ406
XX AAQ406
XX HVR; h
KW HVR; h
KW Animal
KW Chromo
XX Synthe
PN FR2680
PD 26-FEB
XX 22-AUG
PR 22-AUG
PR 22-AUG
PR 22-AUG
PR WPI; 1
PT Peact
PT heredi
PT react
PT polynu
PT react
XX The Se
CC The Se
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             a large polymerase protein, and a partial or complete PIV background genome, or antigenome of a human PIV (HPIV) or bowine PIV (BPIV), combined with one or more heterologous gene(s) or genome segment(s) of a different PIV to form a human-bovine chimeric PIV genome or antigenome. The chimeric PIV is useful in attenuated vaccines to elicit immune response against one or more virus(es) selected from HPIVI, HPIV2 and HPIV3 and most preferably against HPIV3. The present sequence represents a BPIV3 parent nucleotide fragment. This sequence is changed in a recombinant BPIV3 for constructing a chimeric PIV.
The sequence is that of a polynucleotide probe which may be used in the detection of new hypervariable regions (HVR) in a DNA sequence. HVR represent a fingerprint useful in e.g. forensic science, paternity testing, animal breeding, etc. The probe may be used as part of a method for the efficient detection in humans or other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HVR; human; animal; forensic science; paternity testing; diagnosis; animal breeding; hereditary diseases; tumours; allele; loss; chromosomal regions; tumour region identification; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to an isolated human-bovine chimeric parainfluenza virus (PIV) that is infectious and attenuated in humans. The virus comprises a major nucleocapsid protein, a nucleocapsid phosphoprotein,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Isolated human-bovine chimeric parainfluenza virus (PIV), useful in attenuated vaccine to elicits an immune response against one or more virus(es) selected from human PIV1 (HPIV1), HPIV2 and HPIV3 -
                                                                                                                                                                                    Detecting the hypervariable regions of DNA for diagnosing hereditary illnesses and tumours - by hybridising labelled polynucleotides and analysing genomic DNA of individuals whereact with restriction fragments
                                                                                                                                                                                                                                                                                                        WPI; 1993-136548/17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10-AUG-1993 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAQ40606 standard; DNA; 14 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                             22-AUG-1991;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 12 BP; 4 A; 4 C; 1 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                 (ETFR ) ETAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             22-AUG-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 tgttgga 7
|||||||
7 TGTTGGA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4.
                                                                                                                                           Page 13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Page 80; 148pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                 FRANCAIS
                                                                                                                                                                                                                                                                                                                                                                                                                                             91FR-0010516.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             91FR-0010516.
                                                                                                                                           46pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      detection
                                                                                                                                           French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0,:
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Pred. No. 3.3e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      probe 14C12.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
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RESULT 14
AAQ78406
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             Query Match
  Best Local Similarity
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                       The antisense oligonucleotides are useful in the treatment of tumours in which expression of TGF-beta is of relevance for pathogenicity and/or inhibition of pathological angiogenesis. They are used especially for the treatment of the immunosuppressive effect of TGF-beta, augmentation of the proliferation of cytotoxic lymphocytes, treatment of endogenous hyperexpression of TGF-beta,
                                                                                    including glioblastomas, treatment and prophylaxis of skin carcinogenesis, and treatment of oesophageal and gastric carcinomas. See AAQ78352-Q78488. The sequences given in GENESEQ files AAQ78352-Q78407 and AAQ78488 are antisense oligodeoxynucleotides of TGF-beta 1. The sequences given in GENESEQ files AAQ78408-78487 are antisense oligodeoxynucleotides of TGF-beta 2 in the form of
                                                   Sequence
                                                                                                                                                                                                                                                                   Claim 6; Page
                                                                                                                                                                                                                                                                                                     New transforming growth oligo:nucleotide(s) - 1
                                                                            phosphorothioate analogues.
                                                                                                                                                             treatment of breast tumours, neurofibromas and malignant gliomas,
                                                                                                                                                                                                                                                                                                                                           WPI; 1994-358266/44
                                                                                                                                                                                                                                                                                                                                                                   Schlingensiepen R;
                                                                                                                                                                                                                                                                                                                                                                                                                               30-APR-1993;
13-MAY-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Transforming growth factor beta; TGF-beta; antisense; treatment; tumour; anglogenesis; breast tumour; neurofibroma; glioma; glioblastoma; catchiogenesis; carchioma; oesophagus; oesophageal; gastric; gut; immunosuppression; oligonucleotide; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    29-APR-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO9425588-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           animals, without the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10-NOV-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Antisense oligonucleotide hybridising to TGF-beta gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         27 - JUN - 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAQ78406
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAQ78406 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                      (BIOG-) BIOGNOSTIK GES BIOMOLEKULARE DIAGNOSTIK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12 TGTTGGA 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 tgttgga 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   7; Conserv
                                                    14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14 BP; 4 A; 4 C; 2 G; 4 T; 0 other;
                                                   BP;
                                                                                                                                                                                                                                                                                                                                                                               Brysch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                   40;
                                                                                                                                                                                                                                                                                                                                                                                                                              93EP-0107089.
93EP-0107849.
                                                   w
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    94WO-EP01362.
                                                   Α;
                                                                                                                                                                                                                                                                74pp;
                                                                                                                                                                                                                                                                                                                                                                               Ę
100,0%;
                                                  3 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   use of mini-satellites or primary enrichment.
                                                                                                                                                                                                                                                                                                    n factor beta
for treating
                                                                                                                                                                                                                                                                                                                                                                             Schlingensiepen
                                                                                                                                                                                                                                                                  English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        14
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                                                   4
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                                                  G;
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Score 7;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 7; DB Pred. No. 3.3); Mismatches
                                                   4
                                                   Ŧ,
                                                                                                                                                                                                                                                                                              anti:sense
immunosuppression, tumours,
                                                   0
                                                  other;
DB 15;
3.3e+04;
                                                                                                                                                                                                                                                                                                                                                                             G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 14; 1
3.3e+04;
hes 0;
                                                                                                                                                                                                                                                                                                                                                                           Schlingensiepen
                                                                                      files AAQ78408-78487 are 2 in the form of
          Length 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
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Query Match Best Local Matches

Similarity 7; Conserv

Conservative

0;

100.0%;

Score 7; Pred. No. Mismatches

DB 19; 1

Length 14;

0

Gaps

0;

Sequence 14

BP; 4 A; 4 C;

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G; 1 U; 0 other;

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RESULT 1
AAV97198/
ID VARY97198/
XX AAV97199/
XX AAV9
XX OALA
XX Sola
XX Sola
XX WO98
XX WO98
XX WO98
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XX III
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XX OALA
XX WO98
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CC The
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                                                                                                                                                                                                                                                                                                                                alkaloids; or (ii) involved in flower formation. AAV95982 to AAV96334, and AAV96335 to AAV96354 represent potato solanidine glucosyltransferase hammerhead and halirpin ribozymes, respectively. AAV95629 to AAV95981, and AAV96355 to AAV96734 represent potato solanidine glucosyltransferase target sequences. AAV96734 to AAV97170, and AAV97171 to AAV97195 represent potato citrate synthase hammerhead and hairpin ribozymes, respectively. AAV96735 to AAV96772, and AAV97196 to AAV9720 represent potato citrate synthase target sequences. Ribozymes of the present potato citrate synthase target sequences. Ribozymes of the present invention can be used to inhibit the synthesis of toxic alkaloids in invention can be used to inhibit the synthesis of toxic alkaloids.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      24-NOV-1997;
28-JAN-1997;
28-JAN-1997;
                                                                                                                                                solanaceous plants, particularly potato but also tomato, pepper, aubergine and ditura or to inhibit flowering in potato, lettuce, spinach, cabbage, brussel sprouts, arugula, kale, collards, chard, beet, turnip, sweet potato and turf grass. Also the ribozymes can be used for RNA manipulation in the same way that restriction endonucleases are for DNA, as well as to examine genetic drift and mutations in plants and to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention describes enzymatic nucleic acid molecules with RNA-cleaving activity (e.g. ribozymes) which are capable of modulating the expression of plant genes: (1) involved in biosynthesis of alkaloids; or (11) involved in flower formation AAV95982 to AAV96334,
                                                                                             detect specific RNA. The to consensus sequences w
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 54; Page 59; 79pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New enzymatic nucleic acid(s) - useful for, e.g. reducing alkaloid biosynthesis or regulating flowering
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1998-427939/36.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       McSwiggen JA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      14-JAN-1998;
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AAV97198 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      W09832843-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Solanum tuberosum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Potato citrate synthase target sequence position
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              flower formation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Solanidine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Solanidine; glucosyltransferase; potato; citrate synthase; target; nammerhead ribozyme; hairpin ribozyme; alkaloid biosynthesis; flower formation; cleavage; solanaceous plant; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ω
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97US-0036545.
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                                                The ribozymes can be targeted to specific s within a family of related genes, and be present at only very low concentrations.
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Result
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 200000000
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Sequence:
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1: /cgn2_6/ptodata/2,
2: /cgn2_6/ptodata/2,
3: /cgn2_6/ptodata/2,
4: /cgn2_6/ptodata/2,
5: /cgn2_6/ptodata/2,
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/cgn2_6/ptodata/2/ina/5B_COMB.seq:*
/cgn2_6/ptodata/2/ina/6A_COMB.seq:*
/cgn2_6/ptodata/2/ina/6B_COMB.seq:*
/cgn2_6/ptodata/2/ina/PCTUS_COMB.seq:*
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US-09-196-523-25
US-08-303-004-119
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US-08-182-968A-201
US-08-182-968A-201
US-08-363-240A-615
US-08-363-240A-615
US-08-363-240A-616
US-08-774-306A-201
US-08-784-306A-246
US-08-585-684B-1366
US-08-585-684B-1729
US-08-585-684B-1731
US-08-585-684B-1730
US-08-585-684B-1730
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Sequence 13, Appl Sequence 21, Appl Sequence 211, Appl Sequence 246, App Sequence 614, App Sequence 615, App Sequence 616, App Sequence 616, App Sequence 136, App Sequence 1367, App Sequence 1730, App Sequence 1731, App Sequence 1731, App Sequence 1732, App Sequence 1732, App Sequence 1748, App Sequence 1768, App Sequence 216, App Sequence 216, App Sequence 216, App Sequence 216, App Sequence 2176, App Sequence 1176, App Sequen
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Pred. No. 6.7e+03; 0; Mismatches 0; Inde	Score 7; DB 2; Len Pred. No. 6.7e+03; 0; Mismatches 0;	631751A h L. abricated, Flowthrough Porcetus for Discrete Detection ns a Avenue, N.W. ble /MS-DOS se #1.0, Version #1.25 se #1.0, Version #1.25 96 912 HARC0001 0N: 3: Score 7: DB 2: Length ON:	US-09-038-073-1731 US-09-038-073-1768 US-09-038-073-1769 US-09-038-073-1770 US-09-038-073-1770 US-09-038-073-1770 US-08-035-350-32 US-08-166-664-15 US-08-169-461-15 US-07-890-609-15 US-08-072-063-4 US-08-119-773-23 US-08-119-773-23 US-08-281-940-25 US-08-390-850-433 US-08-391-24A-810 US-08-391-124A-810 US-08-441-370-2 US-09-282-146-10
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US-08-303-004-19/c

; Sequence 19, Application US/08303004

; Patent No. 5556955
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OTHER INFORMATION: oligonucleotide
US-09-196-523-25
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Best Local Similarity
Matches 7; Conser
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SEQ ID NO 25
LENGTH: 11
                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/07/931,311B
FILING DATE: 19920818
ATTORNEY/AGENT INFORMATION:
NAME: Berridge, William P.
REGISTRATION NUMBER: 30,024
REFERENCE/DOCKET NUMBER: WPB 28264
TELECOMMUNICATION INFORMATION:
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Patent No. 6248525
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                09-196-523-25/c
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FILE REFERENCE: ILI 130
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Vergnaud, or TITLE OF INVENTION: POTITLE OF INVENTION: POTITICE OF INVENTION: HOUMBER OF SEQUENCES:
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ORGANISM: Artificial Sequence
FEATURE:
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                                                                                                                                                                                                                                                                                                              MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
                                       TELEFAX: (703)
TELEX: 90-1799
                                                                                                                                                                                                                                                        FILING DATE:
CLASSIFICATION: 536
                                                                       TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: Alexandria
STATE: Virginia
                                                                                                                                                                                                                                                                                             APPLICATION NUMBER:
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P.O. Box 19928
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                                836-2787
PTO ALEX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Process for Detection of New Polymor-
phic Loci in an ADN Sequence, Nucleotide Sequences Fo
Hypridisation Probes and Their Biological Applications
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; MOLECULE TYPE: ; DESCRIPTION: US-08-765-340-154
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Patent No.
                                                                                                                                                                                                       TELEPHONE: (212) 758-4
TELEFAX: (212) 751-684
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compati
OPERATING SYSTEM: PC-DOS
SOFTWARE: P1.30 (EPO)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 23-DEC-1996
PRIOR APPLICATION DATA:
APPLICATION UNMBER: JP 145
FILING DATE: 27-JUN-1994
PRIOR APPLICATION UNMBER: JP 311
APPLICATION NUMBER: JP 311
FILING DATE: 21-NOV-1994
ATTORNEY/AGENT INFORMATION:
AMMERICAN SECURITY OF THE PRIOR APPLICATION SECURITY OF THE PRIO
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MOLECULE TYPE:
HYPOTHETICAL: I
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APPLICANT:
                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: 14
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 758-4800
TELEFAX: (212) 751-6849
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                            NAME: SERUNIAN, LESLIE REGISTRATION NUMBER: 35,353
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/08 FILING DATE: 23-DEC-1996
                                                                                           TOPOLOGY:
                                                                                                                 STRANDEDNESS:
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STATE: NEW YORK
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UCHIDA, T.,
TANAKA, Y.,
MATSUDA, Y.,
KONDO, S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           linear
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SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MORGAN & FINNEGAN, L.L.P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   N
O
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                        other nucleic acid
/desc = "synthetic
                                                                                                                 single
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
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6.8e+03
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RESULT 6
US-08-182-968A-246/c
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                                                                                           Sequence 246, Application US/08182968A Patent No. 5610054
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                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
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Patent No. 5610054
                                                                                                                                                                                                                                                                         Matches
GENERAL INFORMATION:
APPLICANT: Draper, Kenneth G.
TITLE OF INVENTION: METHOD AND REAGENT FOR
TITLE OF INVENTION: INHIBITING HEPATITIS C
TITLE OF INVENTION: VIRUS REPLICATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: (213) 955-04-
TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 15
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/08/182,968A
FILING DATE: 13-JANUARY-1994
PRIOR APPLICATION DATA:
APPLICATION UMBER: 07/882,888
FILING DATE: 14-MAY-1992
ATTORNEY/AGENT INFORMATION:
ANALYSIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Draper, Kenneth G.
TITLE OF INVENTION: METHOD AND REAGENT FOR
TITLE OF INVENTION: INHIBITING HEPATITIS C
TITLE OF INVENTION: VIRUS REPLICATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
ZIP: 90071-2066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 20
                                                                                                                                                                                                                                                                                                                                                                            STRANDEDNESS: sir
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE:
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CORRESPONDENCE ADDRESS:
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Query Match
Best Local Similarity
7; Conserve
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US-08-363-240A-614/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION UMBER: US/08/182
FILING DATE: 13-JANUARY-1994
PRIOR APPLICATION NUMBER: 07/882,88
APPLICATION NUMBER: 07/882,88
FILING DATE: 14-MAY-1992
ATTORNEY/AGENT INFORMATION:
NAME: WATBUTG, RICHARD J.
REGISTRATION UMBER: 32,327
REFERENCE/DOCKET NUMBER: 205/
TELECOMMUNICATION INFORMATION:
TELEFAX: (213) 489-1600
TELEFAX: (213) 955-0440
TELEY: 67-3510
                                                                                                                                                                                                                                          APPLICANT: Couture, Larry
APPLICANT: McSwiggen, James
APPLICANT: McSwiggen, Charles
APPLICANT: Bisgaier, Charles
APPLICANT: Pape, Michael
TITLE OF INVENTION: METHOD AND
TITLE OF INVENTION: PROVENSION
TITLE OF INVENTION: PROGRESSION
TITLE OF INVENTION: OF VASCULA
                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
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                 COUNTRY: U.S.A.
ZIP: 90071
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OPERATING SYSTEM: IBM P.C. SOFTWARE: Word Perfect 5.1 CURRENT APPLICATION DATA:
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MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: storage
COMPUTER: IBM Compatible
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                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
                                                                                                             CITY: Los Angeles
STATE: California
                                                                                                                                                ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
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STRANDEDNESS: singl
TOPOLOGY: linear
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STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         13 TGTTGGA
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                                                                                                                                                                                    ADDRESSEE:
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633 West Fifth
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IBM Compatible
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                                                                                                                                                                                                                                                           METHOD AND REAGENT FOR PREVENTION, INHIBITION OF PROGRESSION AND REGRESSION
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OF VASCULAR DISEASES
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Pred. No. 6.8e+03;
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NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 210//
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEPAX: (213) 955-0440
TELEPAX: (213) 955-0440
INFORMATION FOR SEQ ID NO: 614:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
companier services at angle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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Best Local (
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ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Alchard
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 211
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT:
                                                                                                                                                                                                                                             OPERATING SYSTEM: IBM P.C. DOS 5.0 SOFTWARE: Word Perfect 5.1 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/363,240A FILING DATE: December 23, 1994 PRIOR APPLICATION DATA: APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM: TO BE STORY OF THE S
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TITLE OF INVENTION: METHOD AND REAGENT FOR
TITLE OF INVENTION: PROCRESSION AND REGRESSION
TITLE OF INVENTION: OF VASCULAR DISEASES
NUMBER OF SEQUENCES: 1243
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:
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15 TGTTGGA 9
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STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              reaccn 100.0%; Score 7; DB 1; Local Similarity 100.0%; Pred. No. 6.8e+0 res 7; Conservative n. wint.
                                                                                                                                                                                                          FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: U
ZIP: 90071
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STATE: California COUNTRY: U.S.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OPERATING SYSTEM: IBM P.C. DOS 5.0 SOFTWARE: Word Perfect 5.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   McSwiggen, James
Bisgaier, Charles
Pape, Michael
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Couture, Larry
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linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     IBM Compatible
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                                           210/096
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TOPOLOGY:
US-08-363-240A-616
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SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                               INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                            FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: WARDURG, Richard
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 210/
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEFAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Pape, Michael
TITLE OF INVENTION: METHOD AND REAGENT FOR
TITLE OF INVENTION: PROGRESSION AND REGRESSION
TITLE OF INVENTION: OF VASCULAR DISEASES

1743
                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage
COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/08/363,: FILING DATE: December 23, 1994 PRIOR APPLICATION DATA: APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Anno 170
                                                                                                                                                                                                                                                                                                                                             OPERATING SYSTEM: IBM P.C. SOFTWARE: Word Perfect 5.1
                           LENGTH: 15 base pairs TYPE: nucleic acid STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15 TGTTGGA 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity les 7; Conserv
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TELEX: 67-3510
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                               90071
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Bisgaier, Charles
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                 linear
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13) 955-0440
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Pred. No.
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6.8e+03;
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                                                                                         US-08-774-306A-246/c
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                                                   Sequence 246, Application US/08774306A Patent No. 5869253
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                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                    Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/08/774,306A FILING DATE: December 26, 1996 PRIOR APPLICATION DATA: APPLICATION NUMBER: 08/182,968 FILING DATE: January 13, 1994 APPLICATION NUMBER: 07/882,888 FILING DATE: May 14, 1992 ATTORNEY/AGENT INFORMATION: NAME: Warburg, Richard J
GENERAL INFORMATION:
APPLICANT: Draper, Kenneth G.
TITLE OF INVENTION: METHOD AND REAGENT FOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: Storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Draper, Kenneth G.
TITLE OF INVENTION: METHOD AND REAGENT FOR
TITLE OF INVENTION: INHIBITING HEPATITIS C
TITLE OF INVENTION: VIRUS REPLICATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OPERATING SYSTEM: IBM P.C. SOFTWARE: WORD Perfect 5.1 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION: TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
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TELEFAX: (213) 955-0440
TELEX: 67-3510
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Warburg, Richard J. REGISTRATION NUMBER: 32,3 REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: 11
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                                                                                                                                                                                                                                                        100.0%; Score 7; DB 2; 57.1%; Pred. No. 6.8e+03;
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Best Local Similarity
Thes 7; Conserv
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US-08-774-306A-246
                                                                                                                                                                                                                                                                                                                                                                 US-08-585-684B-1366/c
                                                                                                                                                                                                                                                                                                        Sequence 1366, Application US/08585684B Patent No. 5877021 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/774,306A
FILING DATE: December 26, 1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/182,968
FILING DATE: January 13, 1994
APPLICATION NUMBER: 07/882,888
APPLICATION NUMBER: 07/882,888
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
MEDIUM TYPE: storage
                                                                                                                                                                                                 APPLICANT: Jarvis, Thale
APPLICANT: McSwijgen, James
TITLE OF INVENTION: METHOD AND REAGENT FOR THE
TITLE OF INVENTION: INDUCTION OF GRAFT TOLERANCE
TITLE OF INVENTION: AND REVERSAL OF IMMUNE RESPONSES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: May 14, 1992
ATTORNBY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 223/227
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELECOMMUNICATION INFORMATION: TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
 COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
                                                                                                                                                               NUMBER OF SEQUENCES: 2
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                                                                                                                                                                                                                                                                                            APPLICANT:
                                                                    ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: U.S.A. ZIP: 90071-2066
                                     COUNTRY: U
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: (213) 955-0440 TELEX: 67-3510
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VIRUS REPLICATION
                                                                                                                                                                                                                                                                                          Daniel T.
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Pred. No. 6.8e+03
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Indels

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Gaps

MEDIUM TYPE: COMPUTER: II

IBM Compatible storage

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US-08-585-684B-1367/c

; Sequence 1367, Application US/08585684B

; Patent No. 5877021
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US-08-585-684B-1366
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Best Local Similarity 100.
The 7; Conservative
                           COMPUTER: IBM COMPATIBLE
COMPUTER: IBM COMPATIBLE
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: U5/08/585,684B
FILING DATE: January 16, 1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/000,951
FILING DATE: July 7, #995
ATTORNEY/AGENT INFORMATION
NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
REGISTRATION NUMBER: 32,327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Warburg, Richard REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/078
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: 67-3510
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
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SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                     ZIP: 90071
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OPERATING SYSTEM: IBM P.C. DOS 5.0 SOFTWARE: FastSEQ Version 1.5 CURRENT APPLICATION DATA:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/08/585,684B FILING DATE: January 16, 1996 PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/000,951 FILING DATE: July 7, 1995 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: METHOD AND REAGENT FOR THE TITLE OF INVENTION: INDUCTION OF GRAFT TOLERANCE TITLE OF INVENTION: AND REVERSAL OF IMMUNE RESPONSES NUMBER OF SEQUENCES: 2751
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: Lyon & Lyon STREET: 633 West Fifth Street STREET: Suite 4700
                   REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 21
                                                                                                                                                                                                                                                                                                                                                                                        CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                       COUNTRY:
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                 218/078
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; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
US-08-585-684B-1729
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Best Local Similarity 100.
"Thes 7; Conservative
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Query Match 100.0%; Score 7; DB 2; L
Best Local Similarity 100.0%; Pred. No. 6.8e+03;
Matches 7; Conservative 0; Mismatches 0;
                                                                                                                                                                                               REFERENCE/DOCKET NUMBER: 211
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEEX: 67-3510
INFORMATION FOR SEQ ID NO: 1729
                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/585,68

FILING DATE: January 16, 1996

PRIOR APPLICATION DATA:

APPLICATION NUMBER: 60/000,951

FILING DATE: JULy 7, 1995

ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                               SEQUENCE CHARACTERISTICS: LENGTH: 15 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: IBM Compatible
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STRANDEDNESS: single
TOPOLOGY: linear
585-684
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STREET: Safte 4700
CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OPERATING SYSTEM: IBM P.C. DOS
SOFTWARE: FastSEQ Version 1.5
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                                                                                                                                                                                                                                                                                                          NAME: Warburg, Richard
REGISTRATION NUMBER: 3
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X: (213) 955-0440
67-3510
                                                                                                                                                           15 base pairs
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100.0%; Pred. No.
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Indels

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Gaps

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COMPUTER READBLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/585,684B
FILING DATE: January 16, 1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/000,951
FILING DATE: July 7, 1995
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard
REGISTRATION NUMBER: 218/078
TELECOMMUNICATION INFORMATION:
TELEFAX: (213) 489-1600
TELEFAX: (213) 955-0440
TELEFAX: (213) 955-0440
TELEFAX: 15 base pairs
TYPE: Nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
Search completed: December 15, 2001, 03:22:47 Job time: 7562 sec
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US-08-585-684B-1730/c
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Best Local Similarity ivv.
Thes 7; Conservative
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APPLICANT: Jarvis, Thale
APPLICANT: McSwiggen, James
TITLE OF INVENTION: METHOD AND REAGENT FOR THE
TITLE OF INVENTION: INDUCTION OF GRAFT TOLERANCE
TITLE OF INVENTION: AND REVERSAL OF IMMUNE RESPONSES
NUMBER OF SEQUENCES: 2751
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: Suite 4700
CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
TENTERED TOTAL
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13 TGTTGGA 7
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Maximum DB
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Perfect score:
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Listing first 45 summaries
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Copyright (c) 1993 - 2000 Compugen
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16.936 Million cell updates/sec
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BH000596 2M0288N06
AZ361153 1M0104A17
AZ450829 1M0249E13
AZ508369 1M0250A24
AL475173 T. brucei
AZ663995 1M0543M22
AZ819914 2M0091N16
AZ999558 2M0286N24
                                                                                                                   Description
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AUTHORS
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ORGANISM
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KEYWORDS
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                                                                    84112, USA
Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                       Mammalia; Eutheria;
1 (bases 1 to 19)
Dunn, D., Aoyagi, A.,
                Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0145 row: C column: 22
Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
High quality sequence stop: 19.
                                                                                                          Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
                                                                                                                                                             Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
                                                                                                                                                                                                                                                                             AZ386406 19 bp DNA GSS 02-OCT-2000 1M0145C22F Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0145C22 F, DNA sequence.
                                                                                                                                     plasmid inserts
Unpublished (2000)
                                                                                                                                                                                                                               Mus musculus
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AZ386406.1 GI:10500106
                                                                                                                                                      Mouse whole genome scaffolding with paired
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                                                                                                  Rm. 308, Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                          house mouse.
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Rodentia;
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A2587241
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AL475439 T. brucei

AL1074857 oy27bl1.s

AL158948 ui42e07.x

AL1453394 tj37902.x

AL2869367 ZM0181p03

AZ352012 1M0090M13

AZ352012 1M00961N14

AZ419525 T. brucei

AZ777241 ZM011G02

AZ332711 1M0661N14

AZ419519 1M0166N02

AZ3939009 ZM01971.3

AZ580757 1M0369D07

AZ829102 ZM0106K02

AZ942800 ZM0203D12

AL473241 T. brucei

AL473241 T. brucei

AL4333172 1M0062020

AZ357647 1M0099D19

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AZ367843 1M0279F06

AZ320254 1M0429A24

AL476327 T. brucei

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VERSION
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Best Local Similarity
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|||||||
11 TGTTGGA 17
                                                Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0279 row: F column: 05
Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mus musculus

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus

1 (bases 1 to 19)

Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C.,
High quality sequence stop: 19.
Location/Qualifiers
                                                                                                                                                                                           Tel: 801 585 5606 Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                 plasmid inserts
Unpublished (2000)
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                                                                                                                                                                                                                                                                                                    University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                                       Contact: Robert B. Weiss
                                                                                                                                                                                                                                                                                                                                                                                                                              Mouse whole genome scaffolding with paired end reads from 10kb
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ilarity 100.0%;
Conservative
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/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC1M0145C22"
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Pred. No. 3.6
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           Tel: 801 585 5606

Fax: 801 585 7177

Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0
Plate: 0311 row: C column: 24
Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
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A2484701
                                                                                                                                                                                    84112,
                                                                                                                                                                                                                        University of Utah
                                                                                                                                                                                                                                                                     Unpublished (2000)
Contact: Robert B.
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High quality sequence stop: 20
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/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC2M0279F05"
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/note="Vector: PWD42nv; Purified genomic DNA from M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /sex="Female"
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                         Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: (
Plate: 0050 row: G column: 20
Seq primer: CACACAGGAAACAGCTATGACC
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Contact: Robert B. Weiss
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Mouse whole genome scaffolding with paired end reads from 10kb
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
Class: plasmid ends
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,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC1M0311C24"
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Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0288 row: N column: 06
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                  Tel: 801 585 5606 Fax: 801 585 7177
                                                                                                                                              84112, USA
                                                                                                                                                                                          University of Utah University of Utah
                                                                                                                                                                                                                                    Contact: Robert B.
                                                                                                                                                                                                                                                               Unpublished (2000)
                                                                                                                                                                                                                                                                                       plasmid
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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Location/Qualifiers
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/sex="Male"
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/clone="UUGC2M0069G20"
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Pred. No.
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3.6e+05;
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Tel: 801 585 5606
Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Errc
Plate: 0104 row: A column: 17
                                                                                                                                University of Utah Genome Center University of Utah Rm. 308, Biomedical Polymers Res
                                                                                                                                                                                                Unpublished (2000)
Contact: Robert B.
                                                                                                                                                                                                                                                                                      Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D., Weiss,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1M0104A17R Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0104A17 R, DNA sequence.
                                                                                                                                                                                                                                             plasmid inserts
                                                                                                                                                                                                                                                             Mouse whole genome scaffolding with paired end reads from 10kb
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1 (bases 1 to 21)
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High quality sequence stop:
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7; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /db_xref="taxon:10090"
/clone="UUGC2M0288N06"
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/strain="C57BL/6J"
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    h: 10000 Std Error: 0.00 row: A column: 17
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Rodentia;
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Pred. No. 3.6
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                                                                                                                                                                 Unpublished (2000)
Contact: Robert B.
                                    Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                       University of Utah Genome Center University of Utah
                                                                                                                                                                                                          plasmid inserts
                                                                                                                                                                                                                             Mouse whole genome
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Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C.,
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                                                                                                                                                                                                                                                                                                                                                                                                                      house mouse.
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Email: ddunn@genetics.utah.
Insert Length: 10000 Std
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                                                                                                    308,
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                                                                                                    Biomedical
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/strain="C57BL/6J"
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                                                                                                                                                                                                                         scaffolding with paired end reads from 10kb
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                                                                                                 Polymers Research Bldg.,
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Pred. No. 3.6e+05;
; Mismatches 0;
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Error: 0.00
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primer: CACACAGGAAACAGCTATGACC

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Thes 7; Conserv
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                                                                                                             Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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                                                                                                                                                                                                                                                                    Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., ISlam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                 Mus musculus
Email: ddunn@genetics.utah.edu
                                                                                                                                                                                   Unpublished (2000)
                                                                                                                                                                                                        plasmid inserts
                                                                                                                                                                                                                           Mouse whole genome scaffolding with paired
                                                                                                                                                                                                                                                   and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AZ508369.1 GI:10689781
                                                                                                                                                                                                                                                                                                                                                                                                                                         nouse mouse
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                Conservative
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                                                                                        Biomedical Polymers Research Bldg.,
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/clone="UUGC1M0249E13"
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/strain="C57BL/6J"
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Seg primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
                  Constructed at the Institute for Genomic Research (TIGR), Rockville, MD. Genomic DNA isolated from a cloned population of Trypanosoma brucei (TREU927/4 GUTAt 10.1) was mechanically sheared to give a tight size distribution (4 kb). The v + 1 method used for the library construction is
                                                                                                                                                Direct Submission
Submitted (10-DEC-2000) Trypanosoma brucei genome sequencing
project, Sanger Centre, The Wellcome Trust Genome Campus, Hinxton,
Cambridge CB10 1SA, E-mail: barrell@sanger.ac.uk and
nhl@sanger.ac.uk
                                                                                                                                                                                                                                                                             Hall,N., Bowman,S., Lennard,N.J., Doggett,J., Atkin,R., Chillingworth,C., Ormond,D., Harris,B., El-Sayed,N., Hou,L., Melville,S.E., Rajandream,M.A. and Barrell,B.G.
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T. brucei sheared genomic
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described in detail in Smith, H. and Venter, J.C. (Making small
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/note="Vector: PWD42nv; Purified genomic DNA from M.
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/strain="C57BL/6J"
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Details of T. brucei sequencing at the Sanger Centre are available
at http://www.sanger.ac.uk/Projects/T_brucei/.
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AZ663995 23 bp DNA GSS 14-DEC-2000 1M0543M22R Mouse 10kb plasmid UUGC1M library Mus musculus genomic Clone UUGC1M0543M22 R, DNA sequence.
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                                                                                                                                                                                                                                                                                                                                 Tel: 801 585 5606
Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0543 row: M column: 22
Seq primer: CACAAGGAAACAGCTATGACC
Class: plasmid ends
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University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
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Mouse whole genome scaffolding with paired end reads from 10kb
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/strain="TREU927"
/db_xref="taxon:5691"
/clone="175c05"
/lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
//lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
//note="Vector: PWD42nv; Purified genomic DNA from M.
muscullus C57BL/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
//http://www.jax.org/resources/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passage through i
0.005 inch orifice at constant velocity. The sheared DNA
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/clone="UUGC1M0543M22"
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/strain="C57BL/6J"
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2 (bases 1 to 23)
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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2M0091N16R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC2M0091N16 R, DNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Seq primer: CACACAGGAAACAGCTATGACC Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: ddunn@genetics.utah
Insert Length: 10000 Std
Plate: 0091 row: N column
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Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pwDA2 (gil4732114[gb]AF129072.1), a copy-number inducible derivative of plasmid Rl. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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                                                                    /lab_host="E. Coli strain XL10-Gold, T1-resistant, F /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jack Laboratory Mouse DNA Resource
                                                                                                                                                                                                              /organism="Mus musculus"
/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC2M091N16"
/clone_lib="Mouse 10kb plasmid UUGC1M library"
/sex="Male"
(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a
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Dunn,D., Aoyagi,A., Batber,M., Beacorn,T., Duval,B., Hamil,C:,
Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly
,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
                                                                                                                                                                                                                                                                                                                                                                                       Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: (
Plate: 0286 row: N column: 24
Seq primer: CACACAGGAAACAGCTATGACC
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University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Res
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Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                    Class: plasmid ends
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                                             /lab_host="E. coli strain XL10-Gold, T1-resistant, F-"
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (female) was obtained from the Jackson
                                                                                                                                                 /organism="Mus musculus"
/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC2M0286N24"
/clone_lib="Mouse 10kb plasmid UUGC2M library"
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                        Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
(http://www.jax.org/resources/documents/dnares/). The DNA
                                                                                                                                 /sex="Female"
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                                                                                                                                                                                                                                                                      to give a tight size distribution (4 kb). The v + 1 method used for the library construction is described in detail in Smith, H. and Venter, J.C. (Making small insert libraries for whole genome shotgun sequencing projects. Genome Sequencing: A Practical Approach, eds. M. Vaudin and B. Barrell, Oxford University Press, 1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Submitted (10-DEC-2000) Trypanosoma brucei genome sequencing project, Sanger Centre, The Wellcome Trust Genome Campus, Hinxton,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hall.N., Bowman,S., Lennard,N.J., Doggett,J., Atkin,R., Chillingworth,C., Ormond,D., Harris,B., El-Sayed,N., Hou,L., Melville,S.E., Rajandream,M.A. and Barrell,B.G.
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Trypanosoma brucei
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T. brucei sheared genomic DNA clone 143h10,
                                                                                                                                                                                                  Details of T. brucei sequencing at the Sanger Centre are available at http://www.sanger.ac.uk/Projects/T_brucei/.
                                                                                                                                                                                                                                                                                                                                                                                                                           Rockville, MD. Genomic DNA isolated from a cloned population Trypanosoma brucei (TREU927/4 GUTat 10.1) was mechanically sl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                nhl@sanger.ac.uk
Constructed at the Institute for Genomic Research (TIGR),
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cambridge CB10 1SA, E-mail: barrell@sanger.ac.uk and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Direct Submission
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Euglenozoa; Kinetoplastida; Trypanosomatidae;
                                                                                                                                                                                                                                                 Email: nelsayed@tigr.org
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (bases 1 to 24)
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                                                                 /organism="Trypanosoma brucei"
/strain="TREU927"
/db_xref="taxon:5691"
                          /clone="143h10"
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Constructed at the Institute for Genomic Research (TIGR),
Constructed at the Institute for Genomic Research (TIGR),
Rockville, MD. Genomic DNA isolated from a cloned population of
Trypanosoma brucei (TREU927/4 GUTat 10.1) was mechanically shear
to give a tight size distribution (
4 kb). The v + i method used for the library construction is
described in detail in Smith, H. and Venter, J.C. (Making small
insert libraries for whole genome shotgun sequencing projects.
Genome Sequencing: A Practical Approach, eds. M. Vaudin and B.
Barrell, Oxford University Press, 1999).
Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; I
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
                                                                                                                            sequence.
AI074857
                                                                                                                                                          AI074857 25 bp mRNA EST 24-SEP-1998 oy27b11.sl Soares_senescent_fibroblasts_NbHSF Homo sapiens cDNA clone IMAGE:1667037 3' similar to TR:064507 Q64507 SERINE 1 ULTRA HIGH SULFUR PROTEIN. contains PTR5 repetitive element ;, mRNA
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AL475439
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Submitted (10-DEC-2000) Trypanosoma brucei genome sequencing project, Sanger Centre, The Wellcome Trust Genome Campus, Hinxton, Cambridge CB10 1SA, E-mail: barrell@sanger.ac.uk and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hall,N., Bowman,S., Lennard,N.J., Doggett,J., Atkin,R., Chillingworth,C., Ormond,D., Harris,B., El-Sayed,N., Hou,L., Melville,S.E., Rajandream,M.A. and Barrell,B.G.
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Eukaryota; Euglenozoa; Kinetoplastida; Trypanosomatidae;
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                                                                      human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Email: nelsayed@tigr.org
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/clone="206g04"
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/strain="TREU927"
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3.7e+05;
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Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Seq primer: -40ml3 fwd. ET from Amersham
High quality sequence stop: 1.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information. Trace considered overall poor quality Insert Length: 2304 Std Error: 0.00
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NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tumor Gene Index
                                                                                      100.0%;
ilarity 100.0%;
Conservative (
                                                                                                                                                                                                                      double-stranded cDNA was size selected, ligated to Eco RI adapters (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of a modified pT7T3 vector (Pharmacia). Library went through one round of normalization to a Cot = 5. Library constructed by Bento Soares and M.Fatima Bonaldo."
                                                                                                                                                                                                                                                                                                                                                                                                                            /organism="Homo sapiens"
/db_xref="taxon:9606"
/db_xref="taxon:9606"
/clone="IhAGE:1667037"
/clone_lib="Soares_senescent_fibroblasts_NbHSF"
/tissue_type="senescent fibroblast"
/lab_host="H9H10B (amplicillin resistant)"
/lab_host="H9H10B (amplicillin resistant)"
/note="Vector: pT7T3D (pharmacia) with a modified
polylinker V_TYPE: phagemid; Site_1: Not I; Site_2: E;
lst_strand cDNA was primed with a Not I - oligo(dT)
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Copyright (c) 1993 - 2000 Compugen
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Sequence
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             1 (bases 1 to 11)
Newman, P.J. and Santoso, S.S.
Molecular basis of the human platelet bra/brb alloantigen system and applications thereof Patent: US 5516634-A 6 14-MAY-1996;
Location/Qualifiers
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AR120915 Sequence 5
178419 Sequence 21
A88596 Sequence 74
A89563 Sequence 74
A89563 Sequence 74
A8082368 Sequence 74
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AR082368 Sequence 21
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AX007637 Sequence
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                                                                                                                                                Unclassified.

1 (bases 1 to 13)
Draper,K.G., Chowrira,B., McSwiggen,J.
Thompson,J.D.
HIV nef targeted ribozymes
Patent: US 5972704-A 217 26-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  l (bases 1 to 11)
Newman, P. J. and Santoso, S.S.
Molecular basis of the human platelet bra/brb alloantigen system and applications thereof
Patent: US 5516634 A 7 14-MAY-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 7 from patent 120742
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Sequence 217 from patent
178419
178419.1 GI:3014573
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Swiggen,J.A. and Mamone,J.Anthony.
Hammerhead ribozymes for preferred
Patent: US 5646020-A 5 08-JUL-1997;
                                                                                                                                                                                                                                                                                                                                   Unknown.
Unclassified.
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Sequence 5 from patent US
I52063
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Draper,K.G., Chowrira,B., McSwiggen,J.,
Thompson,J.D.
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Brysch, W. and Schilingensiepen, K.
AN ANTISENSE OLIGONUCLECTIDE PREPARATION METHOD Patent: WO 9833904-A 744 06-AUG-1998;
BIOGNOSTIK GES (DE); BRYSCH WOLFGANG (DE)
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Sequence 744 from Patent
A88596
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Patent: US 5693535-A 217 02-F
Location/Qualifiers
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Brysch, W.D. and Schlingensiepen, K.D.
An antisense oligonucleotide preparation method Patent: EP 0856579-A 744 05-AUG-1998;
BIOGNOSTIK GES (DE)
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AR082368
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AR120910
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Draper, K.G., Chowrira, B., McSwiggen, J.,
Thompson, J.D.
Method and reagent for inhibiting human
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HIV nef targeted ribozymes
Patent: US 5972704-A 212 26-OCT-1999;
Location/Qualifiers
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                                     Similarity 7; Conser
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caper,K.G., Chowrira,B., McSwiggen,J.,
                                     100.0%; Score 7; DB 6; Lilarity 100.0%; Pred. No. 5.7e+05; Conservative 0; Mismatches 0;
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212 from patent
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                                                       Length 14;
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Sequence 181 from Patent WO9967428.
AX007639
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Aids-associated retrovirus
Viruses; Retroid viruses; Retroviridae.
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Sequence 212 from
178414
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                                Synthetic
A35660
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                                                                                                                                                                                                                                                                 HIV targeted ribozymes Patent: US 5693535-A 212 02-DEC-1997;
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Draper,K.G., Chowrira,B., McSwiggen,J., Stinchcomb,D.T. and
                                                                                                                                                                                                                                                                                                                                             Unknown.
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artificial sequence.
1 (bases 1 to 15)
Camble,R. and Edge,M.D.
Analogous interferon polypeptides, process
and pharmaceutical compositions containing
Patent: EP 0194006-A 105 10-SEP-1986;
IMPERIAL CHEMICAL INDUSTRIES PLC
                                                                                                                                                                                                                                                                                                                                                         1 (bases 1 to 15) Stinchcomb,D.T., Draper,K., McSwiggen,J. and Jarvis,T. C-myb ribozymes having 2'.5'-linked adenylate residues Patent: US 5817796-A 90 06-OCT-1998; Location/Qualifiers
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                             Hz-1 virus persistence-associated-gene 1 (PAG1) promoter uses therefor, and compositions containing same or products therefrom Patent: US 5911982-A 25 15-JUN-1999;
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                                                                 Chao, 1
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Copyright (c) 1993 - 2000 Compugen
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14.932 Million cell updates/sec
Leptospira nucleot Human dendritic cee Metastatic breast Metastatic breast Metastatic breast teast NORF gene SA Yeast NORF gene SA
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diagnosis;
                       New isolated pathogenic Leptospira bacterium - useful for, edeveloping products for conferring protective immunity, and prophylactic or therapeutic treatment
                                                                                                                                                                                                  (AGRI-)
(PIGR-)
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                                                                                                                 WPI; 1998-520791/44.
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                                                                                                                                                                                                      ) AGRIC VI
) PIG RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            pathogenic Leptospira; protective immunity; therapy;
ss.
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                                                                                                                                                                                                  VICTORIA SERVICES ES & DEV CORP.
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AAF4 1077

AAF4 1323

AAF4 3526

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AAF4 9156

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AAF5 969

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AAX64619
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RSV N hammerhead
Mouse relA hammerh
Human CD40 hammerh
Human CD40 hammerh
Human CD40 hammerh
Human B7-10 hammerh
Human B7-1 hammerh
Human B7-1 hammerh
Human B7-1 hammerh
Human B7-1 hammerh
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HIV-1 protease gen
rb gene antisense
HIV-1 protease gen
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RSV 1C hammerhead
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Claim

15; Page 72; 94pp; English.

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RESULT/
AAZ7857/
ID AAZ787/
XX AAZ78/
AC AAZ7
XX AAZ7
DT 10-A
XX SAGE Huma
XX SAGE KW APC;
XX CYCC
XX CYCC
XX BPR 19---
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                                                                                         19-JUN-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cytotoxic T-lymphocyte; tumour antigen; immunotherapy; anticancer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human dendritic cell SAGE tag,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 19-JUN-1998;
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                                               19-JUN-1998;
19-JUN-1998;
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Local Similarity 100.
nes 7; Conservative
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       98US-0089997

98US-0089997

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98US-00990035

98US-00990040

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98US-00990043

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98US-0089992.
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Pred. No. 1.2
0; Mismatches
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1.2e+08;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           They may be used in vaccines to induce an immune response, particularly care against a tumour antigen; to modulate the genotype of an APC; to screen compared to a sense of the compared to an APC; to screen compared to a symptotist and monitoring of differentially expressed genes in compared to approach these genes. Detection of the dendritic cell cappression of these genes, or of their encoded proteins, can be used to identify cells as belonging to the monocyte lineage. Cells containing these genes can be used in active immunotherapy (or to stimulate production of a population of antigen-specific effector cells) and containing them are used in gene therapy. Co-administration of containing them are used in gene therapy. Co-administration of containing them are used in gene therapy. The succession of the production of a population of antigen-specific effector cells containing them are used in gene therapy. Co-administration of containing the containing them are used in gene therapy. Co-administration of containing the containi
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(ROBE/)
(SHAN/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  insufficient to activate a robust cytotoxic immune response that can lyse the tumour cells, immunostimulatory cofactors also being required for efficient activation of cytotoxic T-lymphocytes (CTLs). Nucleic acid sequences identified using the SAGE tags have several potential uses.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  activation of the cytotoxic immune response, particularly against tumour cells. Tumour antigen presentation via the MHC (major histocompatibility complex) and subsequent recognition by T-cell receptors is alone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           with monocytes. Some of the transcripts correspond to known genes or ESTs (expressed sequence tags) which were previously unknown to be preferentially or differentially expressed in dendritic cells, while other transcripts correspond to novel genes. Antigen-presenting cell (APC) associated costimulatory factors play an important role in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              differentially expressed in monocyte-derived dendritic cells compa
with monocytes. Some of the transcripts correspond to known genes
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08-DEC-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cells, useful in gene vaccines against cancer
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                                                                                                                                                                           AAZ80864;
                                                                                                                                                                                                                          AAZ80864 standard;
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                                                                        Metastatic breast tumour cell upregulated transcript tag #98.
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nilarity 100.
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98US-0111715.
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4.3e+04;
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Human; metastatic breast tumour tissue; breast cancer; non-metastatic breast tumour tissue; gene therapy; anti

anticancer;

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RESULT
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                                                                                                                                                                                                              AZ80767 to AAZ83941 represent tags corresponding to distinct CC transcripts that are preferentially transcribed in the metastatic breast tumour tissue (i.e. are upregulated in metastatic breast tumour cells). CC AAZ8342 to AAZ86677 represent tags corresponding to distinct transcripts CC that are preferentially transcribed in the primary or non-metastatic breast tumour tissue (i.e. are downregulated in metastatic breast tumour cells). These transcripts can be used for diagnosis, prognosis, CC monitoring and treatment of breast cancer, particularly where metastatic. CC Diagnosis is by standard immunoassays or hybridisation/amplification creactions. Compounds that modulate expression of the transcripts are potentially useful for treatment of (metastatic) breast cancer, while cc equences), particularly an antigen-encoding sequence for use in gene or cell-based vaccines. Polypeptides encoded by the transcripts are also useful in vaccines; for diagnosing breast cancer and for raising cspecific antibodies (Ab). Ab are used to detect the polypeptides or as therapeutic agents. Host cells that produce the polypeptides can be used to expand and isolate populations of educated, antigen-specific immune effecter cells, e.g. cytotoxic T lymphocytes, and these used for
                                                                                                                                                                   Query Match
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Metastatic breast tumour cell upregulated transcript
                      07-APR-2000
                                           AAZ82567
                                                             AAZ82567 standard; DNA; 10 BP
                                                                                                                                                                                                             Sequence 10 BP; 3 A; 0 C; 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  prevention and treatment of cancer
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                                                                                                                                                          Conservative
                     (first entry)
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98US-0090041.
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98US-0089997.
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Pred. No. 4.3e+04;
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           AAZ82768;
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                               AAZ82768 standard;
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AZEB0767 to AAZEB3941 represent tags corresponding to distinct CC transcripts that are preferentially transcribed in the metastatic breast tumour tissue (i.e. are upregulated in metastatic breast tumour transcripts CC AAZEB3942 to AAZEB6677 represent tags corresponding to distinct transcripts CC that are preferentially transcribed in the primary or non-metastatic breast tumour cells). These transcripts can be used for diagnosis, prognosis, compounds treatment of breast cancer, particularly where metastatic Diagnosis is by standard immunoassays or hybridisation/amplification creactions. Compounds that modulate expression of the transcripts are potentially useful for treatment of increatatic) breast cancer, while concern that transcripts are used to direct expression, in selected cell types, of e.g. therapeutic genes (also ribozymes or antisense cell types), particularly an antigen-encoding sequence for use in gene or cell-based vaccines; for diagnosing breast cancer and for raising useful in vaccines; for diagnosing breast cancer and for raising concernition and isolate populations of educated, antigen-specific immune effecter cells, e.g. cytotoxic T lymphocytes, and these used for immune center immunorabarance.
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) ROBERTS B L.
) SHANKARA S.
Similarity 7; Conserv
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Conservative
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98US-0089997.
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  AAZ83117
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                                                                                                                                                                                                                                                                                                                                                                                                                                              prevention and treatment of cancer
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                                                                                                                                                        Sequence 10
                                                                                                                                                                              adoptive immunotherapy.
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98US-0089997.
98US-0090039.
98US-0090040.
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tumour tissue (i.e. are upregulated in metastatic breast tumour cells).

AAZ83942 to AAZ86677 represent tags corresponding to distinct transcripts that are preferentially transcribed in the primary or non-metastatic breast tumour tissue (i.e. are downregulated in metastatic breast tumour cells). These transcripts can be used for diagnosis, prognosis, monitoring and treatment of breast cancer, particularly where metastatic. Diagnosis is by standard immunoassays or hybridisation/amplification reactions. Compounds that modulate expression of the transcripts are potentially useful for treatment of (metastatic) breast cancer, while promoters from the transcripts are used to direct expression, in selected cell types, of e.g. therapeutic genes (also ribozymes or antisense sequences), particularly an antisen-encoding sequence for use in gene or cell-based vaccines. Polypeptides encoded by the transcripts are also useful in vaccines; for diagnosing breast cancer and for raising specific antibodies (Ab). Ab are used to detect the polypeptides on as therapeutic agents. Host cells that produce the polypeptides can be used to expand and isolate populations of educated, antigen-specific immune effecter cells, e.g. cytotoxic T lymphocytes, and these used for
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Sequence
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) ROBERTS B
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                                                       immunotherapy.
ВP;
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; vaccine; diagnosis; ss.
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""" hes 7; Conserve
                                                                                                                CC comprising a SAGE (serial analysis of gene expression) tag. Also
CC described are: (1) a method (M1) of using NORF genes to affect the cell
CC cycle comprising administering a NORF gene whose expression varies by at
CC least 10% between any two phases of the cell cycle selected from log
CC antifungal drugs comprising: (a) contacting a test substance with a
CC yeast cell; and (b) monitoring expression of a NORF gene whose
CC expression varies as in M1, where a test substance which modifies the
CC expression of the yeast gene is a candidate antifungal drug; (3) a method
CC (M3) for identifying human genes which are involved in cell cycle
CC progression comprising contacting human DNA with a probe which comprises
CC as in M1; and (4) a method (M4) for identifying a candidate drug as a
CC member of a class of drugs having a characteristic effect on gene
CC expression in a yeast cell comprising contacting a yeast cell with a
CC candidate drug and monitoring expression in the yeast cell of at least 1
CC NORF gene whose expression is affected by the class of drugs. The NORF
CC queses may be used to study, monitor and affect phases of the cell cycle.
CC the differentially expressed genes may be used as markers of phases of
CC the cell cycle. The methods may be used to identify candidate drugs which
CC candidate drugs and monitoring expression in the exemplification of
CC the present invention. AAF33262 to AAF33267 represent linkers and PCR
CC primers used in the SAGE method, in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention describes an isolated DNA molecule comprising coding sequence of a yeast gene selected from a group of 745\ \text{NORF} ()
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Yeast gene coding sequences comprising NORF genes with serial analysis of gene expression (SAGE) tags, useful for studying, monitoring and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              nor previously assigned open reading frame; nonannotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification;
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                                                                                                                Sequence 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             assigned open reading frame; or nonannotated ORF)
                                                                                                                BP;
                 Conservative
                                                                                                              7 A; 1 C; 0 G;
                                   100.0%;
                 0,
                                   Score 7;
Pred. No.
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                                                                                                              2 T; 0 other;
                 Mismatches
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                                     4.3e+04;
                                                       DB 22;
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                 Indels
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                 Gaps
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YEAST: Saccharomyces cerevisiae
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YEAST: Saccharomyces cerevisiae.
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YEAST: Saccharomyces cerevisiae.
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                                                                                                                                                                                                                                                                                                                                                                                Cycle comprising administering a NORF gene whose expression varies by at C least 10% between any two phases of the cell cycle selected from log phase, S phase and G2/M; (2) a method (M2) for screening candidate CC quartifungal drugs comprising; (a) contacting a test substance with a CC yeast cell; and (b) monitoring expression of a NORF gene whose CC expression varies as in M1, where a test substance which modifies the CC expression varies as in M1, where a test substance which modifies the CC expression of the yeast gene is a candidate antifungal drug; (3) a method (CC expression comprising contacting human DNA with a probe which comprises CC at least 10 contiguous nucleotides of a NORF gene whose expression varies at 10 contiguous nucleotides of a NORF gene whose expression varies CC member of a class of drugs having a characteristic effect on gene CC expression in a yeast cell comprising contacting a yeast cell with a CC candidate drug and monitoring expression in the yeast cell of at least 1 CC NORF gene whose expression is affected by the class of drugs. The NORF CC genes may be used to study, monitor and affect phases of the cell cycle. The methods may be used to identify candidate drugs which a fact the cell cycle and for identification of antifuncal drugs which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention describes an isolated DNA molecule comprising a coding sequence of a yeast gene selected from a group of 745 NORF (not previously assigned open reading frame; or nonannotated ORF) genes comprising a SAGE (serial analysis of gene expression) tag. Also described are: (1) a method (M1) of using NORF genes to affect the cell control of the contro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Yeast gene coding sequences comprising of gene expression (SAGE) tags, useful affecting phases of the cell cycle -
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                                                                                                                                                                                                                                              affect the cell cycle and for identification of antifungal drugs. AAF33268 to AAF44064 represent SAGE tags used in the exemplification of the present invention. AAF33262 to AAF33267 represent linkers and PCR
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                                                                                                                                                                                           method,
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                                                                                                                                                                                           in the exemplification of the present
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The present invention describes an isolated DNA molecule comprising a coding sequence of a yeast gene selected from a group of 745 NORF (not previously assigned open reading frame; or nonannotated ORF) genes CC comprising a SAGE (serial analysis of gene expression) tag. Also Cdescribed are: (1) a method (M1) of using NORF genes to affect the cell Cycle comprising administering a NORF gene whose expression varies by at Celleast 10% between any two phases of the cell cycle selected from log CC phase, S phase and G2/M; (2) a method (M2) for screening candidate celligating and (b) monitoring expression of a NORF gene whose cexpression varies as in M1, where a test substance with a CC yeast cell; and (b) monitoring expression of a NORF gene whose cexpression of incomprising contacting a test substance which modifies the cell cycle sexpression varies as in M1, where a test substance which modifies the cexpression varies as in M1, where a test substance which modifies the cell comprising contacting human DNA with a probe which comprises as in M1; and (4) a method (M4) for identifying a candidate drug as a method (M4) for identifying a candidate drug as a candidate drug and monitoring expression in the yeast cell with a CC candidate drug and monitoring expression in the yeast cell of at least 1 CC condidate drug and monitoring expression in the yeast cell of at least 1 CC candidate drug and monitoring expression in the present cycle, the cell cycle and for identification of antifungal drugs. The NORF consenses may be used to study, monitor and affect the cell cycle, which consenses of the cell cycle and for identification of antifungal drugs. CC chaf33268 to AAF44064 represent SAGE tags used in the exemplification of the present invention. AAF33262 to AAF33267 represent linkers and PCR
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AAF38092/
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Best Local Similarity
Matches 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Yeast gene coding sequences comprising of gene expression (SAGE) tags, useful affecting phases of the cell cycle -
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example; Page 172; 419pp; English.
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100.0%; P:
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4.3e+04;
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CC coding sequence of a yeast gene selected from a group of 745 NORF (not Cr previously assigned open reading frame; or nonannotated ORF) genes CC comprising a SAGE (serial analysis of gene expression) tag. Also CC described are; (1) a method (M1) of using NORF genes to affect the cell CC cycle comprising administering a NORF gene whose expression varies by at CC least 10% between any two phases of the cell cycle selected from log CC phase, S phase and G2/M; (2) a method (M2) for screening candidate CC antifungal drugs comprising; (a) contacting a test substance with a CC yeast cell; and (b) monitoring expression of a NORF gene whose CC expression varies as in M1, where a test substance which modifies the CC expression of the yeast gene is a candidate involved in cell cycle and contacting the phase is a candidate and cycle at least 10 contiguous nucleotides of a NORF gene whose expression varies as in M1; where a test substance which comprises CC as in M1; and (4) a method (M4) for identifying a candidate drug as a cCC member of a class of drugs having a characteristic effect on gene CC expression in a yeast cell comprising contacting a yeast cell with a CCC candidate drug and monitoring expression in the yeast cell of at least 1 CCC NORF gene whose expression is affected by the class of drugs. The NORF genes may be used to study, monitor and affect phases of the cell corla
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AAF39829/c
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                                                                                                                                                                                                                                                                                                                                                                                                  Yeast gene coding sequences comprising of gene expression (SAGE) tags, useful affecting phases of the cell cycle -
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                for studying, monitoring
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RESULT 11
AAF40359/c
ID AAF403
XX AAF403
XX AAF403
XX AAF403
XX Yeast;
KW Yeast;
KW Nor pr
KW Serial
KW linker
XX Saccha
XX WO2000
XX 11-JEC
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CC The present invention describes an isolated DNA molecule comprising a CC coding sequence of a yeast gene selected from a group of 745 NORF (not previously assigned open reading frame; or nonannotated ORF) genes CC comprising a SAGE (serial analysis of gene expression) tag. Also CC comprising administering a NORF gene whose expression varies by at CC clast 10% between any two phases of the cell cycle selected from log CC phase, S phase and G2/M; (2) a method (M2) for screening candidate CC antifungal drugs comprising: (a) contacting a test substance with a CC yeast cell; and (b) monitoring expression of a NORF gene whose which modifies the CC expression varies as in M1, where a test substance which modifies the CC expression comprising contacting human pune which a probe which comprises CC at least 10 contiguous nucleotides of a NORF gene whose expression varies as in M1; and (4) a method (M4) for identifying a candidate antifungal drug; (3) a method CC as in M1; and (4) a method (M4) for identifying a candidate antifungal drug; (3) a method CC as in M1; and (4) a method (M4) for identifying a candidate antifungal drug; (3) a method CC as in M1; and (4) a method (M4) for identifying a candidate antifungal drug; (3) a method CC as in M1; and (4) a method (M4) for identifying a candidate antifungal drug; (3) a method CC as in M1; and (4) a method (M4) for identifying a candidate antifungal drug; (3) a method (M4) for identifying a candidate antifungal drug; (3) a method (M5) for identifying a candidate antifungal drug; (3) a method (M5) for identifying a candidate antifungal drug; (3) a method (M5) for identifying a candidate antifungal drug; (3) a method (M5) for identifying a candidate antifungal drug; (3) a method (M5) for identifying a candidate antifungal drug; (3) a method (M5) for identifying a candidate antifungal drug; (3) a method (M5) for identifying a candidate antifungal drug; (3) a method (M5) for identifying a candidate antifungal drug; (3) a method (M5) for identifying a candidate antifungal dru
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Yeast gene coding sequences comprising of gene expression (SAGE) tags, useful
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       nor previously assigned serial analysis of gene
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example; Page 253; 419pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16-JUN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   23-MAR-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 tttgata 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PCR primer; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           phases of the cell cycle
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       gene expression;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  English.
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NORF genes with serial analysis for studying, monitoring and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB
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                                         Matches
                                                                                                                                                                                 member of a class of drugs having a characteristic effect on gene expression in a yeast cell comprising contacting a yeast cell with a candidate drug and monitoring expression in the yeast cell of at least 1 NORF gene whose expression is affected by the class of drugs. The NORF genes may be used to study, monitor and affect phases of the cell cycle, the differentially expressed genes may be used as markers of phases of the cell cycle. The methods may be used to identify candidate drugs which affect the cell cycle and for identification of antifungal drugs. AAF33268 to AAF44064 represent SAGE tags used in the exemplification of the present invention. AAF33262 to AAF33267 represent linkers and PCR primers used in the SAGE method, in the exemplification of the present
                                                                                                                                     Sequence 10
 1 tttgata 7
                                         Similarity 7; Conser
                                                                                                                                     BP;
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RESULT 1
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AAF40389 standard;
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DNA;
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Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF; nor previously assigned open reading frame; nonannotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification; linker; PCR primer; ds.

Yeast NORF 23-MAR-2001

gene SAGE

tag oligonucleotide SEQ ID

NO:7128

(first entry)

Saccharomyces cerevisiae

WO200077214-A2

14-JUN-2000; 2000WO-US16223

16-JUN-1999; 99US-0335032

(UYJO) UNIV JOHNS HOPKINS

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Vogelstein B,

Kinzler K;

WPI; 2001-061874/07.

Yeast gene coding sequences comprising of gene expression (SAGE) tags, useful phases of the cell cycle NORF genes with serial analysis for studying, monitoring and

English

The present invention describes an isolated DNA molecule comprising a coding sequence of a yeast gene selected from a group of 745 NORF (not previously assigned open reading frame; or nonannotated ORF) genes comprising a SAGE (serial analysis of gene expression) tag. Also described are: (1) a method (M1) of using NORF genes to affect the cell cycle comprising administering a NORF gene whose expression varies by at least 10% between any two phases of the cell cycle selected from log phase, S phase and G2/W; (2) a method (M2) for screening candidate antifungal drugs comprising: (a) contacting a test substance with a yeast cell; and (b) monitoring expression of a NORF gene whose varies as in M1, where a test

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RESULT 13
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CC progression comprising contacting human DNA with a probe which comprises
CC at least 10 contiguous nucleotides of a NORF gene whose expression varies
CC as in M1; and (4) a method (M4) for identifying a candidate drug as a
CC member of a class of drugs having a characteristic effect on gene
CC expression in a yeast cell comprising contacting a yeast cell with a
CC candidate drug and monitoring expression in the yeast cell of at least 1
CC NORF gene whose expression is affected by the class of drugs. The NORF
CC genes may be used to study, monitor and affect phases of the cell cycle,
CC the differentially expressed genes may be used as markers of phases of
CC the cell cycle. The methods may be used as markers of phases of
CC affect the cell cycle and for identification of antifungal drugs.
CC AAF33268 to AAF44064 represent SAGE tags used in the exemplification of
CC the present invention. AAF33262 to AAF33267 represent linkers and PCR
CC primers used in the SAGE method, in the exemplification of the present
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The present invention describes an isolated DNA molecule comprising a coding sequence of a yeast gene selected from a group of 745 NORF (not previously assigned open reading frame; or nonannotated ORF) genes comprising a SAGE (serial analysis of gene expression) tag. Also described are: (1) a method (M1) of using NORF genes to affect the cell cycle comprising administering a NORF gene whose expression varies by at
                                                                                                                                                                                                                                                                                                        Yeast gene coding sequences comprising of gene expression (SAGE) tags, useful
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14-JUN-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       serial analysis of gene expression; linker; PCR primer; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF nor previously assigned open reading frame; nonannotated ORF; SAGE;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (UYJO ) UNIV JOHNS HOPKINS.
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ilarity 100.
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Pred. No. 4.3
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            antifungal;
                                                                                                                                                                                                                                                                                                     NORF genes with serial analysis for studying, monitoring and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 22; 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            identification;
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The present invention describes an Example; Page 268; 419pp; English.

isolated DNA molecule comprising

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CCCXXX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CC yeast cell; and (b) monitoring expression of a NORF gene whose CC expression varies as in M1, where a test substance which modifies the CC expression of the yeast gene is a candidate antifungal drug; (3) a method CC (M3) for identifying human genes which are involved in cell cycle progression comprising contacting human DNA with a probe which comprises CC at least 10 contiguous nucleotides of a NORF gene whose expression varies CC as in M1, and (4) a method (M4) for identifying a candidate drug as a CC member of a class of drugs having a characteristic effect on gene CC expression in a yeast cell comprising contacting a yeast cell with a CC candidate drug and monitoring expression in the yeast cell of at least 1 CC NORF gene whose expression is affected by the class of drugs. The NORF CC genes may be used to study, monitor and affect phases of the cell cycle, the differentially expressed genes may be used as markers of phases of the cell cycle. The methods may be used to identify candidate drugs which affect the cell cycle and for identification of antifungal drugs.

CC AAF33268 to AAF44064 represent SAGE tags used in the exemplification of the present invention. AAF33267 to AAF33267 represent linkers and PCR primers used in the SAGE method, in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                               nor previously assigned open reading
serial analysis of gene expression; a
linker; PCR primer; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAF40778 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   least 10% between any two phases of the cell cycle selected from log phase, S phase and G2/M; (2) a method (M2) for screening candidate antifungal drugs comprising; (a) contacting a test substance with a yeast cell; and (b) monitoring expression of a NORF gene whose
                                                                                                                                 WPI; 2001-061874/07.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Yeast NORF gene SAGE tag oligonucleotide SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    23-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAF40778;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 10
                                                                                 of gene expression
                                                                                                                                                                Velculescu V,
                                                                                                                                                                                                                                                                                                                                  WO200077214-A2
                                                                                                                                                                                                                                                                                                                                                                  Saccharomyces cerevisiae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF;
                                                                                              Yeast gene coding sequences comprising
                                                                                                                                                                                               (UYJO ) UNIV JOHNS HOPKINS
                                                                                                                                                                                                                                   16-JUN-1999;
                                                                                                                                                                                                                                                                   14-JUN-2000; 2000WO-US16223
                                                                                                                                                                                                                                                                                                   21-DEC-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    7; Conserv
                                                              phases of the cell cycle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     B₽;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                              Vogelstein B,
                                                                                                                                                                                                                                   9908-0335032
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                                                                                                                                                                                                                                                                                                                                                                                                    gene expression; antifungal;
; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA;
                                                                                 (SAGE) tags,
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Pred. No.
                                                                                                                                                                  Kinzler
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                                                                                 useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                   frame; nonannotated
                                                                               NORF genes with serial analysis for studying, monitoring and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 22;
4.3e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ID NO:7517
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 10;
                                                                                                                                                                                                                                                                                                                                                                                                                  tag; identification;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0
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affecting phases of the cell cycle

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Query Match
Best Local Similarity
~+~hes 7; Conserv:
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AAF41077
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        the cell cycle. The methods may be used to the cell cycle and for identification of antifungal drugs. affect the cell cycle and for identification of antifungal drugs. AAR33268 to AAR44064 represent SAGE tags used in the exemplification of the present invention. AAR33262 to AAR33267 represent linkers and PCR primers used in the SAGE method, in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       previously assigned open reading frame; or nonannotated ORF) genes comprising a SAGE (serial analysis of gene expression) tag. Also described are: (1) a method (M1) of using NORF genes to affect the cell cycle comprising administering a NORF gene whose expression varies by an example of the cell cycle selected from log phase, S phase and G2/M; (2) a method (M2) for screening candidate antifungal drugs comprising: (a) contacting a test substance with a
Yeast gene coding sequences comprising NORF genes with serial analysis of gene expression (SAGE) tags, useful for studying, monitoring and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200077214-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  nor previously assigned open reading frame; nonannotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAF41077;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAF41077 standard; DNA; 10 BP
                                                                                                           WPI; 2001-061874/07
                                                                                                                                                                      Velculescu V,
                                                                                                                                                                                                                                        (UYJO ) UNIV JOHNS HOPKINS.
                                                                                                                                                                                                                                                                                                          16-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                         14-JUN-2000; 2000WO-US16223
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Saccharomyces cerevisiae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Yeast NORF gene SAGE tag oligonucleotide SEQ ID NO:7816
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           23-MAR-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 10
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                                                                                                                                                                   Vogelstein B,
                                                                                                                                                                                                                                                                                                          99US-0335032
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3 A; 1 C;
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100.0%; P
tive 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 7;
Pred. No.
                                                                                                                                                                          Kinzler K;
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The present invention describes an isolated DNA molecule comprising a coding sequence of a yeast gene selected from a group of 745 NORF (not previously assigned open reading frame; or nonannotated ORF) genes comprising a SAGE (serial analysis of gene expression) tag. Also described are: (1) a method (M1) of using NORF genes to affect the cell cycle comprising administering a NORF gene whose expression varies by at least 10% between any two phases of the cell cycle selected from log phase, S phase and G2/M; (2) a method (M2) for screening candidate antifungal drugs comprising: (a) contacting a test substance with a cyeast cell; and (b) monitoring expression of a NORF gene whose expression varies as in M1, where a test substance which modifies the expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for identifying human genes which are involved in cell cycle progression comprising contacting human DNA with a probe which comprises as in M1 and (M) for identifying a candidate antifunce expression varies
                                                                             as in M1; and (4) a method (M4) for identifying a candidate drug as a member of a class of drugs having a characteristic effect on gene expression in a yeast cell comprising contacting a yeast cell with a candidate drug and monitoring expression in the yeast cell of at least 1 NORF gene whose expression is affected by the class of drugs. The NORF genes may be used to study, monitor and affect phases of the cell cycle, the differentially expressed genes may be used as markers of phases of the cell cycle. The methods may be used as markers of phases of the cell cycle and for identification of antifungal drugs. AAF33268 to AAF34064 represent SAGE tags used in the exemplification of the present invention. AAF33262 to AAF33267 represent linkers and PCR primers used in the SAGE method, in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example; Page 279; 419pp; English.
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Sequence 10 BP; 2 A; 2 C; 1 G; 5 T; 0 other

Ş Query Match
Best Local Similarity Matches 1 tttgata Conservative 100.0%; 0,; Score 7; DB 22; Pred. No. 4.3e+04; Mismatches 0; Length 10; Indels 0; Gaps

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Search completed: December 15, Job time: 7884 sec 2001, 03:29:44

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2: /cgn2_6/ptodata/2/ina/5A_COMB.seq:*
3: /cgn2_6/ptodata/2/ina/6A_COMB.seq:*
4: /cgn2_6/ptodata/2/ina/FGTUS_COMB.seq:*
5: /cgn2_6/ptodata/2/ina/PCTUS_COMB.seq:*
6: /cgn2_6/ptodata/2/ina/backfiles1.seq:*
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Gapop 10.0 , Gapext 1.0
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Copyright (c) 1993 - 2000 Compugen Ltd
                                                                                                                                                                                                                                                                                                                                                Length
      DB
1 US-08-086-634-6
1 US-08-086-634-7
4 US-09-042-353-4
4 US-08-758-417A-272
1 US-08-758-417A-272
1 US-08-71-880A-217
2 US-08-910-408-217
3 US-09-249-215-217
2 US-08-910-408-212
2 US-08-910-408-212
2 US-08-91-932A-129
US-08-291-932A-129
US-08-291-932A-130
US-08-291-932A-131
US-08-334-847-9
US-08-334-847-655
US-08-334-847-655
US-08-334-847-656
US-08-334-847-656
US-08-334-847-656
US-08-335-684B-113
US-08-585-684B-113
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            Sequence 6, Appli
Sequence 4, Appli
Sequence 272, App
Sequence 217, App
Sequence 217, App
Sequence 217, App
Sequence 217, App
Sequence 212, App
Sequence 212, App
Sequence 212, App
Sequence 129, Appli
Sequence 130, Appli
Sequence 131, Appli
Sequence 655, App
Sequence 655, App
Sequence 113, Appli
Sequence 113, App
Sequence 113, App
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Sequence 115, App
Sequence 116, App
Sequence 117, App
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tttgata TTTGATA	h S	LE OF INVENTIO LE OF SEQUENC RESPONDENCE AD DORESSEE: POI TREET: P.O. TREET: Wiscons JUNYRY: Madison TATE: Wiscons JUNYRY: MADISON TYPE: FSJ70-1149 PUTER READABLE EDIUM TYPE: IS MOUTER READABLE EDIUM TYPE: IS MOUTER READABLE EDIUM TYPE: WORT REATING SYSTE JOHNEY/AGENT IN AME: SCANION LETERATION NUM PPLICATION NUM PPLICATION REGISTRATION NUM PPLICATION EGISTRATION MAME: SCANION EGISTRATION EGISTRATION EGISTRATION ECOMMUNICATION ELEPHONE: (608) MATION FOR SECUENCE CHARACTEE URNCE CHARACTEE URNCE CHARACTEE ROTH: 11 bas YPE: nucleic TRANDEDNESS: OPOLOGY: 11ne ECULE TYPE: C 6-634-6	4 2 2 2	1111111111111111
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	ty erva	TILE OF INVENTION: Molecula TILE OF INVENTION: Applicat MBER OF SEQUENCES: 12 RRESPONDENCE ADDRESS: ADDRESSE: Foley & Lardner STREET: P.O. Box 1497 CITY: Madison STATE: Wisconsin COUNTRY: USA ZIP: 53701-1497 MEDIUM TYPE: Floppy Disk MEDIUM TYPE: Floppy Disk MEDIUM TYPE: IBM PC compatibl OPERATING SYSTEM: PC-DOS/M SOFTWARE: WORDPER: US/08/ FILING TON NUMBER: US/08/ FILING TON NUMBER: US/08/ FILING TON NUMBER: 30-JUNG-1993 TORNEY/ACENT INFORMATION REGISTRATION NUMBER: 30,13 REFERENCE/DOCKET NUMBER: 30,13 REFERENCE/DOCKET NUMBER: 30,13 STECOMMUNICATION INFORMATION TELEPHONE: (608) 258-4258 TELEPAX: (608) 258-4258 RMATION FOR SEQ ID NO: 6: QUENCE CHARACTERISTICS: LENGTH: 11 base pairs TRANDENESS: double TOPOLOGY: 11 near LECULE TYPE: CDNA to mRNA 86-634-6	ion	000000000000000000000000000000000000000
	100 100 tive	MOLECU Human Applic ESS: & Lardr X 1497 ORM: Compati PC-Dos Compati PC-Dos ATATION IDATA: ID	US/08	16 15 15 15 15 15 15 15 15 15 15 15 15 15
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RESULT 3
US-09-042-353-4
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Best Local Similarity
7; Conserve
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                                                                                                                                                                                                                                                                      Sequence 4, Application US/09042353 Patent No. 6255458
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 7, Applicati
Patent No. 5516634
GENERAL INFORMATION:
                                                                                                                                                                                                                                                       GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: NEWMAN, PE
APPLICANT: SANTOSO, S
TITLE OF INVENTION: I
TITLE OF INVENTION: I
TITLE OF INVENTION: I
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                                                                                                                                                           APPLICANT: Lonberg, Nils
APPLICANT: Kay, Robert M.
APPLICANT: Kay, Robert M.
TITLE OF INVENTION: Transgenic No. 6255458-Human Animals for
TITLE OF INVENTION: Producing Heterologous Antibodies
NUMBER OF SEQUENCES: 421
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: 30,136
REFERENCE/DOCKET NUMBER: 30383/132
REFERENCE/DOCKET NUMBER: 30,236/132
REFERENCE/DOCKET NUMBER: 30,136
REFERENCE/DOCKET NUMBER: 30,136
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WordPortfect, Version 5.1
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy Disk
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                              CORRESPONDENCE ADDRESS:
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LENGTH: 11 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: nucleic acid
STRANDEDNESS: double
                                                    COUNTRY:
                                                                           STATE:
                                                                                                          STREET:
                                                                                                                           ADDRESSEE:
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ZIP: 53701-1497
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STATE: Wiscons
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                                       94111-3834
                                                                                      SSEE: Townsend and Townsend and Crew LLP
T: Two Embarcadero Center, Eighth Floor
San Francisco
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Human Platelet Bra/Brb Alloantigen System and
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Pred. No.
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; MOLECULE TYPE: US-09-042-353-4
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                                                                                                                                                                TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US 08/352,322
FILING DATE: 07-DEC-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/544,404
FILING DATE: 10-OCT-1995
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APPLICATION NUMBER: US,
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APPLICATION NUMBER:
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FILING DATE: 18-NOV-
PRIOR APPLICATION DATA:
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                                             LENGTH: 12 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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OPERATING SYSTEM:
SOFTWARE: Patentl
                                                                                                                                                                                                        NAME: Apple, Randolph REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US 01 FILING DATE: 02-DEC-1996
                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER:
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                                   TOPOLOGY:
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                                                                                                                                     (415)
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SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                       (415) 576-0200
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09-MAR-1994
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26-APR-1993
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10-DEC-1993
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18-NOV-1993
                 DNA (genomic)
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10-OCT-1996
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22-JUL-1993
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; TOPOLOGY: linear NOLECULE TYPE: DNA (genomic); SEQUENCE DESCRIPTION: SEQ ID NO: 272: US-08-758-417A-272
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GENERAL INFORMATION:
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4 TTTGATA 10
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                              REFERENCE/DOCKET NUMBER: 014643-009030US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Lonberg, Nils
                                                                                                                                                               SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                        FILING DATE: 18-NOV-1993
APPLICATION UMBBER: US 08/096,762
FILING DATE: 22-JUL-1993
APPLICATION NUMBER: US 08/053,131
FILING DATE: 26-APR-1993
APPLICATION UMBER: US 07/990,860
FILING DATE: 16-DEC-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/08/758,417A FILING DATE: 02-Dec-1996 CLASSIFICATION: <Unknown>
                                                                                                        LENGTH: 12 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US 08/544,404 FILING DATE: 10-CCT-1995 APPLICATION NUMBER: US 08/352,322 FILING DATE: 07-DBC-1994
                                                                                                                                                                                                                                                                     NAME: Serafini, Andrew T. REGISTRATION NUMBER: 41,303
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US 08/161,739 FILING DATE: 03-DEC-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE: 10-DEC-1993
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APPLICATION NUMBER: US 08/165,699
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                                                                                 RESULT 6
US-08-271-880A-217/c
Sequence 217, Application US/08271880A
Patent No. 5693535
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 5, Application US/08166664 Patent No. 5646020 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                          Query Match
Best Local 9
                                                                        GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
                                  APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Warburg, Richard J. REGISTRATION NUMBER: 32,327 REFERENCE/DOCKET NUMBER: 19 TELECOMMUNICATION INFORMATION: TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: HAMMERHEAD RIBOZYI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: James A. McSwiggen APPLICANT: J. Anthony Mamone
                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
                    APPLICANT:
APPLICANT:
                                                                                                                                                                                                 STRANDEDNESS: SII
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET:
                                                                                                                                                                                                                                                                                        Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 tttgata 7
                                                                                                                                                                                                                                                                       Similarity 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Los Angeles
                                                                                                                                                                                                                                                                                                                                                                                                                 nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                E: Lyon & Lyon
611 West Sixth Street
                                                                                                                                                                                                                                                                     100.0%; ilarity 100.0%; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (213)
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                 Kenneth G. Draper
Bharat Chowrira
James McSwiggen
 Dan T. Stinchcomb
                                                                                                                                                                                                                                                                                                                                                                                              single
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                                                                                                                                                                                                                                                                                          Score 7;
Pred. No.
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                                                                                                                                                                                                                                                                                                         DB 1;
                                                                                                                                                                                                                                                                                            9.1e+03;
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Length 13;

Indels

0;

Gaps

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Query Match
Best Local
                                            COUNTY:
ZIP: 90017
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage
AMPHTER: IBM Compatible
TOMPHTER: IBM P.C. DOS (Version 5.0
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS (Version 5.0)
SOFTWARE: WordPerfect (Version 5.1)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/166,664
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 7;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 4; I
9.1e+03;
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US-08-910-408-217/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local S:
Matches 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 206/
TELECOMMUNICATION: TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEFAX: (213) 955-0440
TELEFAX: (213) 955-0440
INFORMATION FOR SEQ ID NO: 217:
                                                                                                                                                                                                                                                             Sequence 217, Application US/08910408
                                                                                                                                                                                                                                       Patent No. 5972704
                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                            APPLICANT:
APPLICANT:
                                                                      TITLE OF INVENTION: METHOD AND REAGENT FOR INHIBITING TITLE OF INVENTION: HUMÁN IMMUNODEFICIENCY VIRUS TITLE OF INVENTION: REPLICATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
LENGTH: 13 base pairs
TYPE: nucleic acid
                                 NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                   APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OPERATING SYSTEM: IBM P.C. DO SOFTWARE: FastSEQ Version 1.5 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ZIP: 90071

COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/08/271,880A
FILING DATE: July 7, 1994
PRIOR APPLICATION DATA: including application
PRIOR APPLICATION DATA: described below:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: May 14, 1992
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Apport
                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity hes 7; Conserv
                                                                                                                                                                                                                                                                                                                                                        10 TTTGATA 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: 08/103,243 FILING DATE: August 6, 1993 APPLICATION NUMBER: 07/882,886
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: Los Angeles
STATE: California
                   ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                     1 tttgata 7
2: Lyon & Lyon
633 West Fifth Street
                                                                                                                      James McSwiggen
Dan T. Stinchcomb
James D. Thompson
                                                                                                                                                                                   Bharat Chowrira
                                                                                                                                                                                                      Kenneth G. Draper
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         James D. Thompson
VENTION: METHOD AND REAGENT FOR INHIBITING
VENTION: HUMAN IMMUNODEFICIENCY VIRUS
                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Score 7; DB 1; ilarity 100.0%; Pred. No. 9.1e+03 Conservative 0; Mismatches 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 13;
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Best Local Similarity
Thehes 7; Conserve
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                                                                                                                                                                                                                                                                                                                                                                                                                       US-09-249-215-217/c
                                                                                                                                                                                                                                                                                                                                                                    Sequence 217, Applicat Patent No. 6159692 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: 08/271,880
APPLICATION NUMBER: 08/103,243
APPLICATION NUMBER: 08/103,243
APPLICATION NUMBER: 07/882,886
FILING DATE: May 14, 1992
ATTORNEY_AGENT INFORMATION:
NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER: 20
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
08/271,880
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 90071

COMPUTER READABLE FORM:

MEDIUM TYPE: 3.5" Diskette, 1.44 Mb

MEDIUM TYPE: storage

COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: FastSEQ Version 1.5 CURRENT APPLICATION DATA:
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|||||||
10 TTTGATA 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 13 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: U.S.A.
          ZIP: 90071
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
storage
                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
                                                                                                                                                                                                                                                 TITLE OF INVENTION: METHOD AND REAGENT FOR INHIBITING HUMAN IMMUNODEFICIENCY VIRUS
                                                                                                                                                                                                                    NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                     APPLICANT: Kenneth G. Draper
Bharat Chowrira
                                                                                                                 CITY: Los Angeles
STATE: California
     COMPUTER:
OPERATING
                                                                                   COUNTRY: U.S.A. ZIP: 90071
                                                                                                                                                                   STREET: 633 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                                       Application US/09249215
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                                                                                                                                                                                                                                                                                                    James McSwiggen
Dan T. Stinchcomb
                                                                                                                                                  Suite 4700
IBM Compatible SYSTEM: IBM P.C. DOS 5.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%; Score 7; 100.0%; Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US/08/910,408
                                                                                                                                                                                                                    REPLICATION: 232
                                                                                                                                                                                                                                                                                      Thompson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
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RESULT 9
US-08-271-880A-212
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Best Local Similarity 100
Matches 7; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       atent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO: 217: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: James McSwiggen
APPLICANT: Dan T. Stinchcomb'
APPLICANT: James D. Thompson'
TITLE OF INVENTION: METHOD AND REAGENT FOR INHIBITING
TITLE OF INVENTION: HUMAN IMMUNODEFICIENCY VIRUS
TITLE OF INVENTION: REPLICATION
NUMBER OF SEQUENCES: 232
                  APPLICATION NUMBER: US/08/271,880A FILING DATE: July 7, 1994
PRIOR APPLICATION DATA: including ap PRIOR APPLICATION DATA: described be PRIOR APPLICATION DATA: described be
                                                                                                                COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOX
SOCTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Ly
                                                                                                                                                                                                                                                                                           STREET: 633 West F:
STREET: Suite 4700
CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: 08/910,408
FILING DATE: CUNKNOWN>
APPLICATION NUMBER: 08/103,243
FILING DATE: August 6,1993
APPLICATION NUMBER: 07/882,886
FILING DATE: May 14, 1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10 TTTGATA 4
                                                                                                                                                                                                                                                                         COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 tttgata 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REGISTRATION NUMBER: 327
REGISTRATION NUMBER: 327
REFERENCE/DOCKET NUMBER: 206/116
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION:
                                                                                                                                                                                                                                                         90071
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       12, Application US/08271880A 5693535
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 13 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/09/249,215 FILING DATE: 12-Feb-1999
                                                                                                                                                                                                                                                                                             California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEX: 67-3510
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: FastSEQ Version 1.5
                                                                                                                                                                                                                                                                                                                                 633 West Fifth Street
Suite 4700
                                                                                                                                                                                                                                                                         U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bharat Chowrira
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Kenneth G. Draper
                                                                                                                                                                                                                                                                                                                                                                       Lyon & Lyon
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: including appplication
: described below:
   08/103,243
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Pred. No.
                                                                                                                                                                                                                   1.44 Mb
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Best Local Similarity
7; Conserva
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TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 2
SEQUIENCE CHARACTERISTICS:
SEQUIENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Patent No. 5972704
                                          FILING DATE: July 7, 1994
APPLICATION NUMBER: 08/103,243
FILING DATE: August 6, 1993
APPLICATION NUMBER: 07/882,886
FILING DATE: May 14, 1992
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard
NAME: Warburg, Richard
                                                                                                                               FILING DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/271,880
FILING DATE: July 7, 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: August 6, 193
APPLICATION NUMBER: 07/882,886
FILING DATE: May 14, 1992
ATTORNEY/AGENT INFORMATION:
NAME: WarDurg, Richard
                                                                                                                                                                                                                                               COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS
SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: 20
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 206/116
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: James D. Thompson
TITLE OF INVENTION: METHOD AND REAGENT FOR INHIBITING
TITLE OF INVENTION: HUMAN IMMUNODEFICIENCY VIRUS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: James McSwiggen
APPLICANT: Dan T. Stinchcomb
APPLICANT: James D. Thompson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: nucleic acid
STRANDEDNESS: singl
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET: 633 West ru
STREET: Suite 4700
                                                                                                                                                                                                                                APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                          ZIP:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                          90071
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633 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                         U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bharat Chowrira
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Kenneth G. Draper
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            single
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                                                                                                                                                                                                                                  us/08/910,408
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Pred. No.
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9.1e+03;
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TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO:

212:

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; Sequence 212, Application US/09249215
; Patent No. 6159692
; GENERAL INFORMATION:
                                                                                                  APPLICATION NUMBER: 08/910,408
APPLICATION NUMBER: 08/103,243
APPLICATION NUMBER: 08/103,243
APPLICATION NUMBER: 07/882,886
FILING DATE: August 6, 1993
APPLICATION NUMBER: 07/882,886
FILING DATE: May 14, 1992
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 32,327
REJEPHONE: (213) 489-1600
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEFAX: 67-3510
INFORMATION FOR SEQ ID NO: 212:
SEQUENCE CHARACTERISTICS:
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Best Local Similarity
Matches 7; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 14 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 tttgata 7
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COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            James D. Thompson
TITLE OF INVENTION: METHOD AND REAGENT FOR INHIBITING
HUMAN IMMUNODEFICIENCY VIRUS
REPLICATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Kenneth G. Drap
Bharat Chowrira
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/09/249,215 FILING DATE: 12-Feb-1999 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OPERATING SYSTEM: IBM P.C. DOS 5.0 SOFTWARE: FastSEQ Version 1.5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: Los Angeles
STATE: California
                                   LENGTH: 14 base pairs TYPE: nucleic acid STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: U.S.A.
TOPOLOGY: linear NCE DESCRIPTION: SEQ ID NO: 212:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; Score 7; DB 2;
1larity 100.0%; Pred: No. 9.1e+03
Conservative 0; Mismatches
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Dan T. Stinchcomb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               storage
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                                                                                                       US-08-373-124A-90
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Patent No. 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
Matches 7; Conser
    Query Match
Best Local Similarity
Matches 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                              FILING DATE: January 13, 1995
PRIOR APPLICATION DATA:
APPLICATION UMBER: 08/245,466
FILING DATE: May 18, 1994
APPLICATION NUMBER: 08/192,943
FILING DATE: February 7, 1994
APPLICATION NUMBER: 07/987,132
FILING DATE: December 7, 1992
APPLICATION NUMBER: 07/987,422
FILING DATE: AUGUST 26, 1992
APPLICATION NUMBER: 07/986,422
FILING DATE: AUGUST 26, 1992
ATTORNEY/ACENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: McSwiggen, James
APPLICANT: Jarvis, Thale
APPLICANT: Jarvis, Thale
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR
TITLE OF INVENTION: TREATMENT OF RESTENOSIS AND
TITLE OF INVENTION: CANCER USING RIBOZYMES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: 209/035
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
                                                                                                                      TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                              NAME: Warburg, Richard REGISTRATION NUMBER: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3 TTTGATA 9
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                                                                                                                                                                                LENGTH:
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                                                                                                                                                                                                                                           x: (213) 955-0440
67-3510
                                                                                                                                                                                    15 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; ilarity 100.0%; Conservative 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Stinchcomb, Dan T.
100.0%; Score 7; DB 1; I ilarity 100.0%; Pred. No. 9.1e+03; Conservative 0; Mismatches 0.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     UMBER: US/08/373,124A
January 13, 1995
                                                                                                                                                                                                                                                                                                                                              32,327
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Pred. No. 9.1e+03;
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                                             Length 15
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Gaps

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RESULT 14
US-08-291-932A-130
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                                                                                                                                                                   Query Match
Best Local Similarity 42...
Conservative
Sequence 130, Application US/08291932A Patent No. 5658780 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 129, Application US/08291932A Patent No. 5658780
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/08/291,932A
FILING DATE: August 15, 1994
CLASSIFICATION: 514
PRIOR APPLICATION DATA: Including application
PRIOR APPLICATION DATA: described below:
APPLICATION NUMBER: 08/245,466
FILING DATE: May 18, 1994
APPLICATION NUMBER: 07,987;132
FILING DATE: December 7, 1992
ATTORNEY/ACENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                               REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 208,
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: 67-3510
                                                                                                                                                                                                                                                                                                                                                                                             NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: WORD PERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION UNITED TO THE PERFECT STREET STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Draper, ... APPLICANT: McSwiggen, James
                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET: Suite 4700
CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY:
                                                                                                                                                                                                                                                                                                                       STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE:
                                                                                                                                                         1 tttgata 7
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                                                                                                                                                                                                                                                                                                                                          nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Warburg, Richard J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           E: Lyon & Lyon
633 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                         15 base pairs
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                                                                                                                                                                                                                                                                                                          linear
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N: 514
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DISEASES OR CONDITIONS
RELATED TO LEVELS OF
                                                                                                                                                                                                              100.0%; Score 7; [ 42.9%; Pred. No. 9
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                                                                                                                                                                                               4; Mismatches
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Вb
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US-08-291-932A-131
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Best Local Similarity
Matches 3; Conserv
                                                                                                                                              Patent No. 5658780 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:

PRIOR APPLICATION DATA: including application

PRIOR APPLICATION DATA: described below:

APPLICATION NUMBER: 08/245,466

FILING DATE: May 18, 1994

APPLICATION NUMBER: 07/987,132

FILING DATE: December 7, 1992

ATTORNEY/AGENT INFORMATION:

NAME: Warburg, Richard J.

REGISTRATION NUMBER: 32,327

REGISTRATION NUMBER: 32,327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: :
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
                   TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Draper, K
APPLICANT: McSwiggen
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                      APPLICANT: Stinchcomb, Dan T. APPLICANT: Draper, Kenneth G. APPLICANT: McSwiggen, James
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OPERATING SYSTEM: IBM P.C. SOFTWARE: WORD Perfect 5.1 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Anna?
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/08/291,932A FILING DATE: August 15, 1994 CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: U.S.A. ZIP: 90071-2066
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STATE: California
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               RIBOZYME TREATMENT OF
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4; Mismatches
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ADDRESSEE: Lyon & Lyon
STREET: Suite 4700
CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
ZIP: 90071-72066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: World Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/291,932A
FILING DATE: AUGUST 15, 1994
CLASSIFICATION DATA: including application
PRIOR APPLICATION DATA: described below:
APPLICATION NUMBER: 08/245,466
FILING DATE: May 18, 1994
APPLICATION NUMBER: 07/987,132
FILING DATE: May 18, 1994
APPLICATION NUMBER: 07/987,132
FILING DATE: December 7, 1992
ATTORREY/ACENT INFORMATION:
NAME: WAIDLITY, RICHARD 3.327
REFERENCE/DOCKET NUMBER: 208/157
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                              Search completed: December 15, 2001, 03:22:47 Job time: 7562 sec
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Result
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Gapop 10.0 , Gapext 1.0
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Copyright (c) 1993 - 2000 Comp
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A2323820 1M0045A01
A2819886 2M0091H16
A2977536 2M0253B04
A2329387 1M0053C07
A2502952 1M0343C04
A2828775 2M0106B06
A2345476 1M0080P11
A2789315 2M0136M22
A2840208 2M0136H12
A2840208 2M0159N22
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 AL466870
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KEYWORDS
SOURCE
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AZ641145/c
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AUTHORS
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                                                                          84112, USA
Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                    Mus musculus
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AZ641145.1
                                                                                                           Rm.
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ALIGNMENTS

source Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pec, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von and Wright, D., Weiss, R. AZ641145 19 bp 1M0503F04R Mouse 10kb r clone UUGC1M0503F04 R, Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0503 row: F column: 04
Seg primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
Location/Qualifiers 1 (bases 1 to 19) Dunn, D., Aoyagi, A., Eukaryota; Metazoa; Mammalia; Eutheria; University of Utah Genome Mouse whole genome scaffolding with paired end Biomedical Polymers Research Bldg., GI:11764838 Weiss Chordata; Rodentia; plasmid UUGClM , DNA sequence. Center Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus. library Mus musculus genomic Duval,B., 20 S. 2030 Pedersen,T., Reilly on Niederhausern,A. reads from 10kb 14-DEC-2000 Hamil,C., en,T., Reilly Ħ SLC, TD

/clone="UUGC1M0503F04" /organism="Mus musculus" /strain="C57BL/6J" /db_xref="taxon:10090"

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REFERENCE
AUTHORS
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AZ323820/c
FEATURES
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                                             Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: |
Plate: 0045 row: A column: 01
Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.

1 (bases I to 20)

1 (bases I to 20)

1 (bases I to 20)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
                                                                                                                                                                                                                                                                                              Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
                                                                                                                                                                                     Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                 Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                plasmid inserts
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                         High quality sequence stop: 20
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                                                                                                                                                                                                                                                                          Biomedical Polymers Research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /sex="Male"
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100.0%; Pred. No. 4.2e+05;
Live 0; Mismatches 0;
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Contact: Robert B. W
University of Utah G
University of Utah
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AZ819886 20 bp DNA GSS AV-FED AVERD 2M0091H16R Mouse 10kb plasmid UUGCIM library Mus musculus genomic clone UUGC2M0091H16 R, DNA sequence.
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                                          Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.
Plate: 0091 row: H column: 16
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sclurognathi; Murldae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GSS
High quality sequence stop: 20
                     Class: plasmid ends
                                                                                                                                     Tel: 801 585 5606 Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                        plasmid inserts
                                                                                                                                                                                                                                                                                                                                             Mouse whole genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   house mouse.
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/strain="C57BL/6J"
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/note="Vector: PWD42nv; Purified genomic DNA from M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /sex="Male"
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Pred. No.
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                        Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0253 row: B column: 04
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                        University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Res
                                                                                                                                                                                                                                                                                                                                                         Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Roye, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.,
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2M0253B04R Mouse 10kb plasmid UUGC2M library Mus clone UUGC2M0253B04 R, DNA sequence.
AZ977536
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mus musculus
Class: plasmid ends
                                                                                                                                                                                                                                                             Contact: Robert B.
                                                                                                                                                                                                                                                                                    Unpublished (2000)
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                                                                                                                      801 585 5606
801 585 7177
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                                                                                                                                                                                        Biomedical Polymers Research Bldg.,
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/clone_lib="Mouse 10kb plasmid UUGClM library"
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Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0053 row: C column: 07
Seq primer: CACACAGGAAACAGCTATGACC
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Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,G., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
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Location/Qualifiers
                                                                                            Tel: 801 585 5606 Fax: 801 585 7177
                                                                                                                                                                                                                                                                         Mouse whole genome scaffolding with paired end plasmid inserts
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and Wright,D.,Weiss,R.
                                                                                                                                                                                 University of Utah University of Utah
                                                                                                                                                                                                                            Contact: Robert B
                                                                                                                                                                                                                                                 Unpublished (2000)
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/strain="C57BL/6J"
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                                                                                                                                                                      Mouse whole genome scaffolding with paired end reads from 10kb plasmid inserts Unpublished (2000) Contact: Robert B. Weiss University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                       Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AZ502952 22 bp DNA GSS 1M0342C04R Mouse 10kb plasmid UUGC1M library Mus clone UUGC1M0342C04 R, DNA sequence.
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
Plate: 0342 row: C column: 04
                                                                          Tel: 801 585 5606
Fax: 801 585 7177
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/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC1M0053C07"
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     column: 04
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Pred. No. 4.2e+05;
0; Mismatches 0;
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Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A., and Wright,D.,Weiss,R.
                                                                                                                                 Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AZ828775 22 bp DNA GSS 20-FEB-2001 2M0106B06F Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC2M0106B06 F, DNA sequence.
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                                             Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                       Mouse whole genome
                                                                                                                                                                                                                                                                                                                                                                                                                                          Mus musculus
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
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/note="Vector: PWD42nv; Purified genomic DNA from M.
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/clone="UUGC1M0342C04"
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Pred. No.
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                                                                                                                                                                                                                          AZ345476 23 bp DNA GSS 29-SEP-2000 1M0080P11F Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0080P11 F, DNA sequence.
                                                                                                 Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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Seq primer: CGTTGTAAAACGACGGCCAGT
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Mammalia; Eutheria;
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                                                                                                                                                                Unpublished (2000)
                                                                                                                                                                                                     Mouse whole genome scaffolding with paired
                                                                                                                                                                                                                                                                                                                                                                                                                                        AZ345476.1
Email: ddunn@genetics.utah.edu
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                 801 585 5606
801 585 7177
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/clone="UUGC2M0106B06"
/clone_1ib="Mouse 10kb plasmid UUGC1M library"
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/note="Vector; PWD42nv; Purified genomic DNA from M.
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/strain="C57BL/6J"
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                                                                                                                                                                                                                                                                                                                                       Mus musculus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                         Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
                                                                                                                                                                                                                                 Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A., and Wright,D.,Weiss,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AZ789315 23 bp DNA GSS 16-FEB-2001
2M0036M22R Mouse 10kb plasmid UUGCIM library Mus musculus genomic
clone UUGC2M0036M22 R, DNA sequence.
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                                               84112,
                                                                                                                                                                 Unpublished (2000)
                                                                                                                                                                                          plasmid inserts
                                                                                                                                                                                                                Mouse whole genome
                                                                                                                                                                                                                                                                                                                                  Mammalia; Eutheria;
1 (bases 1 to 23)
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                                                                     308,
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                                                                  Biomedical
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/note="Vector: PMD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
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/clone="UUGC1M0080P11"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GI:12929986
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100.0%; Pred. No. 4.2e+05;
Live 0; Mismatches 0;
                                                                                                                                                                                                             scaffolding with paired end reads from 10kb
                                                                     Polymers Research Bldg.,
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AUTHORS
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Best Local Similarity
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                                                                                                    plasmid inserts
Unpublished (2000)
Contact: Robert B.
                                                                                                                                                                                         Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                         Mus musculus

Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalla; Butheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus

1 (bases 1 to 23)
                                                             University of Utah Genome Center University of Utah
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                                                                                                                                                                       Mouse whole genome
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Location/Qualifiers
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Plate: 0036 r
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Insert Length: 10000 Std Erro
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                                         308,
801 585 5606
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (http://www.lax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-completent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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                                  Biomedical Polymers Research Bldg.,
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/note="Vector: PWD42nv; Purified genomic DNA from M.
/noscelus C57BL/6J (male) was obtained from the Jackson
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/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC2M0036M22"
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Pred. No. 4.2
0; Mismatches
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4.2e+05;
hes 0;
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RESULT 11
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Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
                                                                                                                                                            Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Musinae; Mus 1 (bases 1 to 23)

Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Welss, R.
                                                                                                                                                                                                                                                                                                                                                                                                                        AZ862305 23 bp DNA 2M0169N22F Mouse 10kb plasmid UUGC1M library Mus musculus clone UUGC2M0169N22 F, DNA sequence.
                                                                                                    Unpublished (2000)
                                                                                                                               plasmid inserts
                                                                                                                                                Mouse whole genome scaffolding with paired
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Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: (
Plate: 0136 row: H column: 12
Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to
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/note="Vector: PWD42nv; Purified genomic DNA from M.
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/clone="UUGC2M0136H12"
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/strain="C57BL/6J"
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RESULT 12
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7; Conserva
nhl@sanger.ac.uk
Constructed at the Institute for Genomic Research (TIGR),
Rockville, MD. Genomic DNA isolated from a cloned population of
Trypanosoma brucei (TREU927/4 GUTat 10.1) was mechanically sheared
                                                                                        Direct Submission
Submitted (10-DEC-2000) Trypanosoma brucei genome sequencing project, Sanger Centre, The Wellcome Trust Genome Campus, Hicambridge CB10 1SA, Etmail: barrell@sanger.ac.uk and
                                                                                                                                                                                                                                                                                                                                                                                                                                          genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TAl144D04Q 23 bp DNA GSS
T. brucei sheared genomic DNA clone 144d04,
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Class: plasmid ends
High quality sequence stop: 23.
Location/Qualifiers
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Insert Length: 10000 Std Error: 0.00
Plate: 0169 row: N Column: 22
                                                                                                                                                                                             Hall.N., Bowman,S., Lennard,N.J., Doggett,J., Atkin,R., Chillingworth,C., Ormond,D., Harris,B., El-Sayed,N., Hou,L., Melville,S.E., Rajandream,M.A. and Barrell,B.G.
                                                                                                                                                                                                                                                                                                                                          Trypanosoma brucei
                                                                                                                                                                                                                                                                                                                                                                                                                           AL466870.1 GI:11836225
                                                                                                                                                                                                                                                                                                                    Eukaryota; Euglenozoa; Kinetoplastida; Trypanosomatidae;
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/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
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/clone="UUGC2M0169N22"
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11 TTTGATA 17
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
1 (bases 1 to 24)
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Details of T. brucel sequencing at the Sanger Centre are available at http://www.sanger.ac.uk/Projects/T_brucel/.
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4 kb). The v + i method used for the library construction is described in detail in Smith, H. and Venter, J.C. (Making small insert libraries for whole genome shotgun sequencing projects. Genome Sequencing: A Practical Approach, eds. M. Vaudin and B. Barrell, Oxford University Press, 1999).
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                                                                                                                                                                                                                                                                            Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0134 row: M column: 14
Seg primer: CGTTGTAAAACGACGGCCAGT
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University of Utah Genome Center
University of Utah
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Fax: 801 585 7177
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                /lab_host="E. Coli strain XL10-Gold, T1-resistant, F
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jack
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /organism="Trypanosoma
/strain="TREU927"
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/clone="UUGC1M0134M14"
Laboratory Mouse DNA Resource
                                                                                                /clone_lib="Mouse 10kb plasmid UUGC1M library"
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Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostom Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; P. 1 (bases 1 to 24)

Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reil,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern, and Wright,D., Weiss,R.
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Insert Length: 10000 Std Error: 0.00
Plate: 0302 row: P column: 17
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/lab_host="E. Coli strain XL10-Gold, Tl-resistant, F
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jack
                                                                                                                 /organism="Mus musculus"
/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="TUGC1M0302P17"
                                                                                          /clone_lib="Mouse 10kb plasmid UUGC1M library"
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                                                                       /sex="Male"
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                                                                                                                                                                                                                                                                                                                                                                           Tel: 801 585 5606
Fax: 801 585 7177
Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.0
Plate: 0543 row: E column: 19
Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AZ663958 24 bp DNA GSS
1M0543E19R Mouse 10kb plasmid UUGC1M library Mus
clone UUGC1M0543E19 R, DNA sequence.
AZ663958
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Islam,H., Longacre,
,M., Rose,M., Rose,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                      High quality sequence stop: 24.
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    and Wright, D., Weiss, R.
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                                                                                                                                      /organism="Mus musculus"
/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC1M0543E19"
                                                                     /clone_lib="Mouse 10kb plasmid UUGC1M library"
/sex="Male"
/lab_host="E. Coli strain XL10-Gold, T1-resistant, F
/note="Vector: PWD42nv; Purified genomic DNA from M.
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Sciurognathi; Muridae; Murinae; Mus.
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Query Match 100.0%; Score 7; DB 13; Length 24; Best Local Similarity 100.0%; Pred. No. 4.2e+05; Matches 7; Conservative 0; Mismatches 0; Indels
1 tttgata 7
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24 TTTGATA 18
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Search completed: December 15, 2001, 02:33:44
Job time: 4984 sec

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

REFERENCE AUTHORS TITLE JOURNAL MEDLINE REFERENCE	LOCUS DEFINITION ACCESSION VERSION KEYWORDS SEGMENT SOURCE ORGANISM	RESULT 1	C 45		C 41	- ω	ωω	u w	0 0 3 3 4 1 4 5	ωι	c c 30	29	c 27	c 25	24	งผ	NN	c 19	ىر د		c 14	<u>_</u>		_	8 ~	امر	C 4	3 2 P	Result
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rimates; Catarrhini L., Conrad,D.G. and ation by lovastatin lecular cloning and alene synthase 7), 12818-12824 (19)	DNA PF synthase gene, intro 6 6 hordata; Craniata; Ve	ALIGNMENTS	AC090544	AC014851 AC087450	AC083839	AC090570 AE002828	AC090590 AF271897	AC090996	CEC34B4 HSABLGR2		CEM03B6 AY027893S6	AC006805	AE002102	AC014767 AP000145	AE006627	MMTHRECO2)45	ω,	2	AX148805 A45819	SPGROELGN	LFU60594 DDU66913	AX148806		BTAJ9224 BTAJ9223		SRSS	HSSSG03 HUM4STS161 G50458	ID
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Direct Submission
Submitted (02-MAR-1998) Biochemistry and Molecular Biology,
Uniformed Services University of the Health Sciences, 4301 Jones
Bridge Road, Bethesda, MD 20814-4799, USA
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                  PCR components: 25 ng of human genomic DNA, 10 pmol of each oligonucleotide, 200 micro-M dNTPS, 0.25 U Taq polymerase (Cetus) in 10 micro-l of 50 mM KCl-20 mM Tris-HGl, pH 8.3 (at room temp), 2.5 mM MgCl-2. Initial denaturation at 94degC for 1.5 min, then 30 cycles of 94degC for 15 sec, 62degC for 3 sec, and 72degC for 3 sec, followed by a final extension at 72degC for 3.5 min, using a perkin-Elmer 9600 thermocycler. PCR-amplified product size 279 bp. sequence submitted by:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 435)

Goold,R.D., disibio,G., Xu,H., Lang,D.B., Dadgar,J., Magrane,G., Dugaiczyk,A., Smith,K.A., Cox,D.R., Masters,S.B. and Myers,R.M. The development of sequence-tagged sites for human chromosome 4 Hum. Mol. Genet. 2 (8), 1271-1288 (1993)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STS; human chromosome 4; sequence tagged site.
Homo sapiens, clone C4-177 from Los Alamos National Laboratory
chromosome 4 cosmid library, plate 4-1-2R, pos3C; vector SuperCos 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human chromosome
                                                                                                                                                                                                       University of Californ'ia San Francisco
San Francisco, CA 94143-0925 USA
Phone: (415) 502-1612 Fax: (415) 476-8391
e-mail: hymcprobes@cgl.ucsf.edu.
Location/Qualifiers
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1 (bases 1 to 668)
Olivier, M. and Cox, D.R.
Unpublished, Olivier, M., C
Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     G50458 668 bp SHCC-79916 Human Homo s G50458 G50458.1 GI:5221635 STS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tel: (650) 320-5800
Fax: (650) 320-5801
Email: olivier@shgc.stanford.edu
Primer A: GCACATPAGCTCTTGGATTGCTT
Primer B: TGCCTTGTGGTTTAGCAATTTTT
STS size: 278
                                                                                                                                                                                                       BAC
and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: Michael Olivier, David R. Cox
Stanford Human Genome Center
Stanford University. School of Medicine
4005 Miranda Ave. 2nd Fl., Palo Alto, CA 94025, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
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KCl:
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                                                                                        complement(286. .30
                                                                                                                                                /map="8"
                                                                                                                                                            /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                       /clone_lib="Human"
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                                  63.6%;
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Best Local Similarity 63.6
Matches 14; Conservative
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Small round structured virus genomic RNA,
containing ORF2 and ORF3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Direct Submission
Submitted (17-CCT-1994) to the DDBJ/EMBL/GenBank databases. Etsuko
Submitted, National Institute of Health; 1-23-1 Toyama,
Shinjuku-ku, Tokyo 162, Japan (Tel:03-5285-1111(ex.2562),
Fax:03-5285-1177)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2 (bases 1 to 1015)
Utagawa, E.T., Takeda, N., Inouye, S., Kasuga, K. and Yamazaki, S. 3'-terminal sequence of a small round structured virus (SRSV)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cDNA to genomic RNA. small round structured virus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Viruses; ssRNA positive-strand viruses, Caliciviridae; Norwalk-like viruses.
                   AJ009225
AJ009225.1
                                                       Bos taurus
                                                                        BTAJ9225
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                                                                                                                                                                                                                                                                                                                                                     Protein_id="BAA07550.1"
/db_xref="GI:123213"
/db_xref="GI:123213"
/td=xref="GI:123213"
/translation="maja1IGAIAASAAGSALGAGIQAGAEAALQAQRYQQDLTLQQN
/translation="maja1IGAIAKNLNTRYSLLQAGGLSSSDAARAVAGAPVTRLVDWGG
SFNHDKEMLGYQMEMSNKLLAKNLNTRYSLLQAGGLSSSDAARAVAGAPVTRLVDWGG
VRVAARQSSATTLRSGNFMAVPLFAQKKPLASEGYSNPAYDPVQRTASWVQSQNSS
RSWSYYHRQALQTYWNTPPGSTSSSSSYSTAPRGYFNTDRLPLFANLRR"
a 251 c 233 g 302 t
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/db_xref="G1:123112"
/tanslation="YPCLLPOEYITHEISEQAPIQGEAALLHYVDPDTNRNLGEEKLY
PGGYLTCVPNSSSTGPQQLPLDGVFVFASWVSRFYQLKPVGTAGPARGRLGVRR"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note="SRSVs are currently recognised as
calicivirus."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /organism="small round structured virus"
/isolate="patient; SRSV-CHIBA-407/87/J"
/db_xref="taxon:37141"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note="ORF3"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /note="ORF2"
                                                         mariner
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                                                                                                                                    Submitted (04-JUN-1998) Bigot Y., Institut de Recherche sur la Biologie de l'Insects, UPRESA CNRS 6035, Facult des Sciences, l Grandmont, 37200 Tours, FRANCE
                                                                                                                                                                                                                                                       Meunier-Rotival,M. and Bigot,Y. Features of the mammal mari transposons and mouse genomes and implications for t Mamm. Genome 11 (12), 1111-1116 (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HSAJ9229 1257 bp DNA PRI 06-JAN-2001 Homo sapiens mariner related transposon Hsmarl, clone mlehsc3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Submitted (04-JUN-1998) Bigot Y., Institut de Recherche sur la Biologie de l'Insects, UPRESA CNRS 6035, Facult des Sciences, Parc Grandmont, 37200 Tours, FRANCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Demattei,M.V., Auge Gouillou,C., Pollet,N., Hamelin,M.H., Meunier-Rotival,M. and Bigot,Y.

Features of the mammal mar1 transposons in the human, she and mouse genomes and implications for their evolution Mamm. Genome 11 (12), 1111-1116 (2000)
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                                                                                                                                                                                                                                                                                                                                    Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 1257)
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                                                                                                                                                                                        Direct Submission
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                                                                                                                                                                                                                                                                                                                           Demattei, M.V., Auge-Gouillou, C., Pollet, N.,
                                                                                                                                                                                                                                                                                                                                                                                                                   human
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                 381
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                 b
            /organism="Homo sapiens"
/transposon="Hsmar1"
/db_xref="taxon:9606"
/clone="mlahsc3"
a 263 c 264 g 349
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/transposon="Hsmarl"
/db_xref="taxon:9913"
/clone="btmlec4"
a 263 c 263 g 34
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Pred. No. 1.1e+03;
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their evolution
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Query Match Best Local Similarity

63.6%;

Score Pred.

14; No.

DB 9; 1.1e+03;

Length 1257;

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BASE COUNT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Demattel, M.V., Auge-Gouillou, C., Pollet, N., Hamelin, M.H., Meunier-Rotival, M. and Bigot, Y. Features of the mammal mari transposons in the human, sheep, and mouse genomes and implications for their evolution Mamm. Genome 11 (12), 1111-1116 (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Submitted (04-JUN-1998) Bigot Y., 1
Biologie de l'Insects, UPRESA CNRS
Grandmont, 37200 Tours, FRANCE
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1 (bases 1 to 1258)
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AJ009224.1
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                                                    Demattei,M.V., Auge-Gouillou,C., Pollet,N., Hamelin,M.H., Meunier-Rotival,M. and Bigot,Y. Meunier-Rotival,M. and Bigot,Y. Reatures of the mammal mari transposons in the human, sheep, and mouse genomes and implications for their evolution Mamm. Genome 11 (12), 1111-1116 (2000)
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AJ009223.1 GI:9187449
             Bigot, Y.
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1 (bases 1 to 1276)
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                           (bases 1 to 1276)
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/transposon="Hsmar1"
/db_xref="taxon:9913"
/clone="btmlec3"
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S 6035, Facult des Sciences,
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Biologie de l'Insects, UPRESA CNRS 6035, Facult des Sciences, I
Grandmont, 37200 Tours, FRANCE
Location/Qualifiers
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14; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Direct Submission
Submitted (21-MAR-1996) Hugh M. Robertson, Entomology, University
of Illinois at Urbana-Champaign, 505 S. Goodwin, Urbana, IL 61801,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Gene 205 (1-2), 203-217 (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   USA
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Robertson, H.M. and Zumpano, K.L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    human
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Robertson, H.M.
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nilarity 63.6%;
Conservative
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/transposon="Hsmar1"
/db_xref="taxon:9913"
/clone="btmlec2"
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1. .1287
                                                                                                                                                             /translation="memmldkkqiraifleeekmgrkaaettrninnaegpgtanert vqwapkkeckgoesledeersgrpsevdndqlraiieadplittrevaeelnvdhstvvrhlkqigkvkkldkwvphelsenqknrreevassliirnnnepeldrivtodekwil ydnrrepaqwldreeapkhepkpnlhqkkvmvtvwsaaglihyselnpgetitseky ydnrrepaqwldreeapkhepkpnlhqkkvmvtvwsaaglihyselnpgetitseky aqqiidemhrklqrapalvnrkgpillhdnarphvaqptlqklnelgyevlphppysp
                                                                                                                                                                                                                                                  /product="mariner transposase"
/protein_id="AAC52010.1"
/db_xref="GI:1263081"
                                                                                                                                                                                                                                                                                                                   /rpt_type=inverted
179. .1210
                                                                                                                                                                                                                                                                                                                                                                  /note="consensus sequence based on 20 unique sequences"
                                                                                   /rpt_type=inverted
284 c 278 g
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                                                                                                                                                                                                                                                                                                                                                                                 /transposon="Hsmarl"
                                                                                                                                                                                                                                                                                                                                                                                                   /organism="Homo sapiens"
/db_xref="taxon:9606"
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   63.6%;
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1.1e+03;
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I (bases 1 to 1306)

Wright, J.A., Young, A.H. and Dugourd, Martisense Oligonucleotide sequences inhibitors of microorganisms
Patent: WO 0136625-A 8 25-MAY-2001, GeneSense Technologies Inc. (CA)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Streptococcus pyogenes.
Streptococcus pyogenes
Bacteria; Firmicutes; Bacillus/Clostridium
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Streptococcus pyogenes
Bacteria; Firmicutes; Bacillus/Clostridium
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Wright, J.A., Young, A.H.
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/db_xref="taxon:1314"
235 c 308 g 356 t
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/db_xref="taxon:1314"
238 c 312 g 349 t
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Pred. No. 1.1e+03;
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Pred. No. 1.1e+03;
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2 (bases
                                                                                                                                                                       Submitted (15-AUG-1996) Dept. of Biology 0322, University of California, San Diego, 9500 Gilman Drive, La Jolla, CA 92093, USA
                                                                                                                                                                                                                                                     Dictyostelium discoideum Dictyostelium discoideum
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U66913
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LFU60594 1481 bp
Leptospira fainei 16S
                                                                                                                                                           California, San Diego, 9500
Location/Qualifiers
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Billinghurst,M.L., Letocart,M., Merien,F.,
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Adler, B., Chappel, R.J.,
                                                                                                                                                                                                   Direct Submission
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                                                   /cell_line="AX4"
join(370..732,813.
/note="ORF DG1040"
<370..732
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /organism="Leptospira fainei"
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join(370. .732,813. .>2321)
/note="ORF DG1040; initially
disrupted gene"
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335 c 439 g 314 t
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                                                                                                         /db_xref="taxon:44689"
                                                                                                                    /organism="Dictyostelium discoideum"
/strain="AX4"
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Pred. No. 1.1e+03;
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                plasmid
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Streptococcus pyogenes.
Streptococcus pyogenes
Bacteria; Firmicutes; Bacillus/Clostridium group; Streptococcaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Direct Submission
Submitted (29-JUN-1995) A. Podbielski, Institut f
Mikrobiologie, an der Rhein Westf. Techn. Hochschu
Pauwelsstr Klinikum, 52057 Aachen, FRG
2 (bases 1 to 2683)
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Podbielski, A.
                                                                                                                                                                                                                                                                                                                                                                                                                     Unpublished
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LISNLTKKSYKSNSSELSQIINQYGSQAEIYIFRCLUVDSIDFKNTNTFITTSFTTSAF
TSTTTSTFTTPSTTASSTAASSSSUQLKIQLFKDEFSRLTKQPHFISVLCRAFDYF
TSTTTSTFTPSTASSTAASSSSUQLKIQLFKDEFSRLTKQPHFISVLCRAFDYF
TSTTTSTFTPSTAASSTAASSSSUQLKIQLFKDEFSRLTKQPHFISVLCRAFDYF
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TNNNNNNNNENNDENEKTKIPYLTFNDLNDKETSNTSPTSSSPKQQSENESSSTTTTT
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/db_xref="G1:2462692"
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/isolate="CS101"
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Streptococcus pyogenes.
Streptococcus pyogenes
Streptococcus pyogenes
Bacteria; Firmicutes; Bacillus/Clostridium group; Streptococcaceae;
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c 22	13	9	1079	20	AAX20676	Polynucleotide seq
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	13	9	1669	22	AAC90031	SEQ ID 1. Staphyl
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c 35	13	9.	3519	21	AAZ94941	Human carbohydrate
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	13	9	4012	18	AAV74358	Staphylococcus aur
38	13	59.1	4254	22	AAI61065	Human polynucleoti
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c 40	13	59.1	4803	22	AAD11574	Human phospholipas
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C 42	13	59.1	4875	22	AAD11386	Human phospholipas
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                                                                                                                Leptospira sp.
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New isolated pathogenic Leptospira bacterium - useful for, e. developing products for conferring protective immunity, and f prophylactic or therapeutic treatment

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                                                                                L. fainei. The LS bacteria can be used for conferring protective immunity against pathogenic LS bacteria in humans or animals. The bacteria can also be used for prophylactic or therapeutic treatment of infections. The DNAs and antibodies may also be used for detection and
                                                                                                                                                                                                         New isolated pathogenic Leptospira bacterium - useful for, e.g developing products for conferring protective immunity, and for prophylactic or therapeutic treatment
                                               Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                     Chappel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Infection; pathogenic Leptospira; protective immunity; therapy; diagnosis; ss.
                     12-OCT-2001
                                                                                                                                                                                                                                                                                                                                   Claim 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         07-MAR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             06-MAR-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAV58900;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAV58900 standard;
                                         AAI15869;
                                                              AAI15869 standard;
                                                                                                                                                                                                               Sequence 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17-SEP-1998.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Leptospira sp
                                                                                                                                                                                                                                                                                                                                                                                                                                        (AGRI-) AGRIC VICTORIA SERVICES (PIGR-) PIG RES & DEV CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO9840099-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Leptospira nucleotide
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                                                                                                                            tgttggannnnnnnntttgata
                                                                                                                  tgttggannnnnnnntttgata
                                                                                                                                                                                                                                                                                                                                                                                                1998-520791/44.
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                                                                                                                                                                                                                                                                                                                                  Page 72; 94pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                               BP; 3 A; 0 C; 4 G; 7 T; 8 other;
                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                    (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         97AU-0005494
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             98WO-AU00145
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA;
                                                              DNA;
                    entry)
                                                                                                                                                                    63.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             sequence
                                                              500
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                                                                                                                                                                      Score 14;
Pred. No.
                                                                                                                                                             Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     DIY LID
                                                                                                                                                                       DB 19;
77;
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                                                                                                                                                            0;
                                                                                                                                                                                 Length
                                                                                                                                                             Indels
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Probe #5802

for gene expression

analysis

'n human

cervical cell sample

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X P X P X O X X X X E X T X C X I D
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                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
Matches 14; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     30-JUN-2000;
03-AUG-2000;
21-SEP-2000;
                                                                                                                                                                                                                                                                                                                                           The present invention relates to human single exon nucleic acid probes (SENP). The present sequence is one such probe. The SENPs are derived from human HeLa cells. The SENPs can be used to produce a single exon microarray, which can be used for measuring human gene expression in a sample derived from human cervical epithelial cells. By measuring gene expression, the probes are therefore useful in grading and/or staging of diseases of the cervix, notably cervical cancer.

Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     04-FEB-2000;
26-MAY-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 25;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         09-AUG-2001.
                                                                     genetic disorder;
                                                                                                                                                                                                                                                                                                                Sequence 500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Penn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    30-JAN-2001; 2001WO-US00670
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO200157278-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cervical cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Probe; human; microarray;
         09-AUG-2001
                             WO200157272-A2
                                                                               Probe; microarray; human;
                                                                                                  Probe #6460 used to measure gene expression in human placenta
                                                                                                                         17-OCT-2001
                                                                                                                                                                AAI37774 standard; DNA; 500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (MOLE-)
                                                                                                                                                                                                                     65
                                                                                                                                                                                                                              1 tgttggannnnnnnntttgata
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                                                                                                                                                                                                                                                                                                                                    ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SG,
                                                                                                                                                                                                                    tgttggaattttccatttgata
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      genome-derived single exon nucleic acid probes useful for zing gene expression in human cervical epithelial cells -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULAR DYNAMICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQ ID No 5802; 487pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hanzel DK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2000US-0180312.
2000US-0207456.
2000US-0608408.
2000US-0632366.
2000US-0234687.
2000US-0234685.
2000US-02346359.
                                                                                                                                                                                                                                                              Conservative
                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                BP; 171 A; 80 C; 92 G;
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                                                                      ss.
                                                                                                                                                                                                                                                                      63.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Chen W,
                                                                               placenta; antenatal diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 gene
                                                                                                                                                                                                                    86
                                                                                                                                                                                                                                        22
                                                                                                                                                                                                                                                             0
                                                                                                                                                                                                                                                          Score 14; DB 22; L
Pred. No. 1.1e+02;
Pred. No. 1.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Rank
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 expression; cervical epithelial cell;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   English.
                                                                                                                                                                                                                                                                                                                157 T; 0 other;
                                                                                                                                                                                                                                                                                Length 500;
                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                             0;
                                                                                                    sample
                                                                                                                                                                                                                                                            Gaps
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AAX99627/c
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Best Local S
Matches 14
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26 - MAY - 2000;
30 - JUN - 2000;
03 - AUG - 2000;
21 - SEP - 2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention relates to single exon nucleic acid probes (SENP). The present sequence is one such probe. The probes are useful for producing a microcarray for predicting, measuring and displaying gene expression in samples derived from human placenta. The probes are useful for antenatal diagnosis of human genetic disorders.
Detection of Ureaplasma urealyticum using novel genes, probes primers % \left( \frac{1}{2}\right) =\frac{1}{2}\left( \frac{1}{2}\right) ^{2}
                                                                                                                                                                                                                                                                                              Ureaplasma urealyticum; nucleic acid detection; infection; pathogen; human urogenital tract; pregnancy; neonatal disease; drug therapy;
                                                                                                                                                                                                                                                                                                                                                                    05-OCT-1999
                                                                                                                                                                                                                                                                                                                                                                                                                          AAX99627 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-488897/53.
                                                                      Cassell GH,
Lefkowitz E;
                                                                                                                                                                                                  05-AUG-1999
                                                                                                                                                                                                                                                      Ureaplasma urealyticum.
                                                                                                                                                                                                                                                                                 suppurative arthritis;
                                                                                                                                                                                                                                                                                                                                          Nucleic acid sequence from U. urealyticum.
                                                                                                                                                                                                                                                                                                                                                                                                 AAX99627;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human genome-derived single exon nucleic acid probes useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            04-OCT-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            27-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30-JAN-2001;
                                           WPI; 1999-469343/39
                                                                                                                                          30-JAN-1998;
                                                                                                                                                                      29-JAN-1999;
                                                                                                                                                                                                                              WO9939007-A1.
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                                                                                                                (UABR-) UAB
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14; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQ ID No 6460; 654pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                gene expression in human placenta
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                                                                                                                RES FOUND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           B₽;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2001WO-US00663
                                                                                    Chen EY,
                                                                                                                                                                                                                                                                                                                                                                    (first entry)
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2000US-0236359
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                                                                                                                                            98US-0073189
                                                                                                                                                                     99WO-US01972.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   63.6%;
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                                                                                    Glass JI,
                                                                                                                                                                                                                                                                                     ss.
                                                                                                                                                                                                                                                                                                                                                                                                                             972
                                                                                                                                                                                                                                                                                                                                                                                                                            ВP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 14; DB 22; | Pred. No. 1.1e+02; 0; Mismatches 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Rank DR;
                                                                                    Glass
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           157 T; 0 other;
                                                                                    JS,
                                                                                      Heiner
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 500;
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Claim

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Page

82; 110pp; English.

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RESULT 7
AAH56865/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention provides methods for the detection and diagnosis CC of Ureaplasma urealyticum infection. It provides novel genes CC (AAX99501-681) that can be used as a source of primers and probes for the CC detection and/or quantification of U. urealyticum in a biological CC sample. The probes that can be used in the method of the invention by CC forming target:probe complex is complementary to a region selected from CC one of the 181 nucleic acid sequences (AAX99501-681). U. urealyticum is CC an opportunistic pathogen of the human urogenital tract that is a CC significant cause of adverse pregnancy outcome, neonatal disease, and CC suppurative arthritis. As the infections are commonly asymptomatic, it is important to have specific and sensitive methods for detecting their CC presence in a patient. Also, as the pathogen has no current antibiotic CC directed specifically against it, it would be advantageous to isolate and CC detect gene sequences which are unique to it, and utilise these as a CC basis for diagnosis of U. urealyticum infection as well as to develop new CC and improved drug therapies. The present invention provides such novel CC polynucleotide sequences (AAX99501-681).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Antisense oligonucleotide; groE; groEL; groES; inhibitor; growth; microorganism; Escherichia coli; Streptococcus pneumoniae; diagnosis; Streptococcus progenes; Staphylococcus aureus; Pseudomonas aeruginosa; antibacterial; antiviral; antiproliferative; antisense therapy;
                                                                                       Novel antisense compounds targeting nucleic acid encoding groEL or groeS gene of microorganism, which hybridize with and inhibit expression of the genes, useful to inhibit growth of microorganism having the genes -
   The present invention specifically claims AAH56368 antisense oligonucleotides to nucleotide sequences
                                                       Disclosure; Fig 11; 110pp; English.
                                                                                                                                                                                  WPI; 2001-355633/37.
                                                                                                                                                                                                                    Wright JA,
                                                                                                                                                                                                                                                                                                                                                                                                    WO200136625-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                      Streptococcus pyogenes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         microbial infection; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Antibiotic resistant S. pyogenes groEL gene partial sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   06-SEP-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAH56865;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAH56865 standard; DNA; 1305
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                          18-NOV-1999;
                                                                                                                                                                                                                                                                                                                              20-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               931 TGTTGGACTATTAGGTTTGATA 910
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                                                                                                                                                                                                                                                       GENESENSE TECHNOLOGIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         972 BP;
                                                                                                                                                                                                                    Young AH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                              2000WO-CA01347
                                                                                                                                                                                                                                                                                        9905-0166249
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     63.6%;
                                                                                                                                                                                                                  Dugourd
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 14; DB 20; L
Pred. No. 1.2e+02;
Pred. No. 1.2e+02;
                                                                                                                                                                                                                    D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         98 G; 438 T; 0 other;
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to AAH568
encoding
                    AAH56832 which
 groE.
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RESULT 8
AAH56862/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CC generally, antisense compounds (I) comprising antisense oligonucleotides CC of 5-50 bases targeted to a nucleotide sequence encoding groEL (heat CS shock protein (HSP)0) (GL) and groEE (HSP10) (GS) gene from a CC GS of a microorganism, where the antisense compound is complementary to GL or CC GS of a microorganism and specifically hybridises with and inhibits the CC expression of GL or GS, is claimed. (I) have antibacterial, antiviral CC and for inhibition of expression of groEE or groEL. (I) are useful for cinhibiting expression of GL or GS in cells or tissues in vitro. (I) are useful for inhibiting the growth of a microorganism, or inhibiting the expression of GL or GS gene in a microorganism (a bacterial cell or CC also useful for treating a gene which involves administering to the microorganism or to a cell infected with the microorganism, (I). (I) are CC also useful for treating a mammalian pathological condition mediated by the microorganism shich involves identifying a eukaryotic organism cor is inhibited. The antisense compounds are utilised for diagnostics, to prevent or delay microbial infections in humans. They are also useful as molecular weight markers. AAH56362 to AAH56367 and AAH56870 represent CC examplification of the present invention. AAH56870 represent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                      microorganism; Escherichia coli; Streptococcus pneumoniae; diagnosis; Streptococcus pyogenes; Staphylococcus aureus; Pseudomonas aeruginosa; antibacterial; antiviral; antiproliferative; antisense therapy;
           Novel antisense compounds targeting nucleic acid encoding groEL or groES gene of microorganism, which hybridize with and inhibit expression of the genes, useful to inhibit growth of microorganism
                                                                                                                                                                                                                                                                                                                                                                                                                           Antisense oligonucleotide; groE; groEL; groES; microorganism; Escherichia coli; Streptococcus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Antibiotic resistant S. pyogenes groEL gene partial sequence SEQ ID:8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    06-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAH56862;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAH56862 standard; DNA; 1306
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          exemplification of the present invention. AAH56855 to AAgroe{o} nucleotide sequence given in the present invention.
                                                                                                                                                                                                                                                                                                     WO200136625-A2
                                                                                                                                                                                                                                                                                                                                    Streptococcus pyogenes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 1305 BP; 406 A; 238
                                                                                   WPI; 2001-355633/37
                                                                                                                                                                                           18-NOV-1999;
                                                                                                                                                                                                                              20-NOV-2000; 2000WO-CA01347.
                                                                                                                                                                                                                                                               25-MAY-2001
                                                                                                                                                        (GENE-) GENESENSE TECHNOLOGIES INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                661 TGTTGGAGCTCCTACTTTGATA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 tgttggannnnnnnntttgata 22
                                                                                                                    JA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                         infection; ds.
                                                                                                                   Young AH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first
                                                                                                                                                                                           9905-0166249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     63.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                640
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 14;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ç
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                                                                                                                      Ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          312 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 22; LC
. 1.3e+02;
. 2.3 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            349 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                            inhibitor; growth;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 1305;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
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RESULT
AAV58896
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Best Local S
Matches 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      also useful for treating a mammalian pathological condition mediated by the microorganisms which involves identifying a eukaryotic organism having a pathological condition mediated by microorganisms having a GL or GS gene and administering (I) such that the growth of microorganism is inhibited. The antisense compounds are utilised for diagnostics, therapeutics, prophylaxis and as research reagents and kits, e.g., to prevent or delay microbial infections in humans. They are also useful as molecular weight markers. AAH56362 to AAH56357 and AAH56833 to AAH56854 represent PCR primers for groE sequences which are useful.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       and antiproliferative activities, and can be used in antisense therapy and for inhibition of expression of groES or groEL. (I) are useful for inhibiting expression of GL or GS in cells or tissues in vitro. (I) are also useful for inhibiting the growth of a microorganism, or inhibiting the expression of GL or GS gene in a microorganism (a bacterial cell or a virus) having a GL or GS gene which involves administering to the microorganism or to a cell infected with the microorganism. (I). (I) are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        generally, antisense compounds (I) comprising antisense oligonucleotides of 5-50 bases targeted to a nucleotide sequence encoding groEL (heat shock protein (HSP)60) (GL) and groES (HSP10) (GS) gene from a microorganism, where the antisense compound is complementary to GL or GS of a microorganism and specifically hybridises with and inhibits the expression of GL or GS, is claimed. (I) have antibacterial, antiviral
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  represent PCR primers for groE sequences which are used in the exemplification of the present invention. AAH56855 to AAH56870 groE nucleotide sequence given in the present invention.
                                                                                                   (AGRI-)
(PIGR-)
                                                                                                                                                   07-MAR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAV58896 standard; DNA; 1477 BP.
   New isolated pathogenic Leptospira bacterium
                                    WPI; 1998-520791/44.
                                                                                                                                                                                   06-MAR-1998;
                                                                                                                                                                                                                      17-SEP-1998.
                                                                                                                                                                                                                                                     WO9840099-A1
                                                                                                                                                                                                                                                                                    Leptospira fainei
                                                                                                                                                                                                                                                                                                                       diagnosis;
                                                                                                                                                                                                                                                                                                                                       Infection;
                                                                                                                                                                                                                                                                                                                                                                                                         20-JAN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                          AAV58896
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 1306 BP; 407 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               antisense oligonucleotides to nucleotide sequences encoding groE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           667
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 tgttggannnnnnnntttgata
                                                                                                                                                                                                                                                                                                                                                                        fainei nucleotide sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TGTTGGAGCTCCTACTTTGATA 646
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        l Similarity
14; Conserv
                                                                                                   AGRIC VICTORIA SERVICES PIG RES & DEV CORP.
                                                                                                                                                                                                                                                                                                                     pathogenic
ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                   97AU-0005494
                                                                                                                                                                                    98WO-AU00145
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                specifically claims AAH56368
                                                                                                                                                                                                                                                                                                                                     Leptospira; protective immunity; therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        235 C; 308 G;
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                                                                                                                    PTY
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1.3e+02;
8;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0 other;
   useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 1306;
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   for,
e.g
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This sequence represents a Leptospira DNA sequence isolated from the pathogenic Leptospira (LS) bacterium of the invention. The bacterium belongs to serogroup Hurstbridge or serovar hurstbridge or the species L. fainei. The LS bacteria can be used for conferring protective immunity against pathogenic LS bacteria in humans or animals. The bacteria can also be used for prophylactic or therapeutic treatment of infections. The DNAs and antibodies may also be used for detection and
                                                                             Human mariner transposase 19 and encoded polynucleotide, applicable diagnosis and treatment of cancer, haemopathy, HIV infection, immunological diseases and inflammation -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             developing products for conferring protective immunity, and prophylactic or therapeutic treatment
                                                         Claim
                                                                                                                             WPI; 2001-418030/44.
P-PSDB; AAG64431.
                                                                                                                                                                                                                                                                                     WO200146240-A1
                                                                                                                                                                                                                                                                                                                                                                                                               Human; mariner transposase 19; anti-HIV; immunostimulatory; cytostatic; antiinflammatory; diagnosis; treatment; cancer; haemopathy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human mariner transposase 19 encoding cDNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAH47810 standard; cDNA; 1573
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 1477 BP; 390 A; 334 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               diagnosis of past or present LS infection.
                                                                                                                                                                                       (BIOW-)
                                                                                                                                                                                                              22-DEC-1999;
                                                                                                                                                                                                                                      18-DEC-2000;
                                                                                                                                                                                                                                                                                                                                                                                                    immunological disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  154
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14; Conserv
                                                                                                                                                                                       BIOWINDOW GENE DEV INC SHANGHAI
                                                         Page 30-31;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Page 69-70;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                      2000WO-CN00601
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                              99CN-0125674.
                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers 1026..1535
                                                                                                                                                                                                                                                                                                           /product= "mariner transposase 19"
/note= "CDS claimed in claim 6"
                                                                                                                                                                                                                                                                                                                             /*tag= a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          63.6%;
                                                        40pp;
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                                                         Chinese
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score
Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         439 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            14;
No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1
1.3e+02;
8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          314 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
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The present sequence is that of the human mariner transposase 19 encode CDNA with anti-HIV, immunostimulatory, cytostatic and antiinflammatory activity. The polypeptide and encoded polynucleotide are applicable in the diagnosis and treatment of cancer, haemopathy, HIV infection.

19 encoding

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AAX86155/c
ID AAX86155 standard; DNA; 1661 BP
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Matches
                                                                                                         Query Match
                                                          Matches
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                                                                                                                                                                                                                             immunosuppressive so provide an increased response to any conjugated or fused antigen. Also, where used for cancer control, they lack the side effects associated with endotoxins. They can also be used to detect specific antibodies and in treatment or prevention of tumours (e.g. sarcoma or cancers of breast, ovary, prostate, lung, pancreas or liver). The Hap60 polynucleotide is used for recombinant production of the protein, as a source of primers and probes for detecting streptococci in standard hybridization/amplification assays, and therapeutically in gene therapy vectors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present sequence encodes a heat shock protein, designated Hsp60-2. The protein, its fragments, variants and fusion proteins, are used to elicit or enhance an immune response against Streptococcus, and to elicit a similar response to a target antigen fused to the protein. Unlike other immunological carriers, Hsp60 proteins are not the protein.
                                                                                                                                                                                   Sequence 1661 BP; 512 A; 308 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 3; Fig 4A-B; 176pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New nucleic acid encoding heat shock protein-60 from Streptococcus, useful in vaccines, as carriers for other immunogens, as anticancer agents and for diagnosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     31-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Streptococcus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Heat shock protein; Hsp60-2; immune response; cancer control; tumour; sarcoma; cancer; gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA encoding a Streptococcus pyogenes heat shock protein (Hsp)60-2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAX86155;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          P-PSDB; AAY23904.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1999-430397/36.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mizzen L, Wisniewski J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (STRE-) STRESSGEN BIOTECHNOLOGIES CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    29-DEC-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   W09935270-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            22-SEP-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      immunological diseases and inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         415
                                                                               Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity 63.0 les 14; Conservative
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     tgttggannnnnnnntttgata
                                                          14;
                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     pyogenes
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                               Score 14;
Pred. No.
                                                                                                                                                                                   407
                                                          Mismatches
                                                                                                                                                                                   <u>ი</u>
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.ches 8;
                                                                               1.3e+02;
                                                                                                                                                                                   434 T; 0 other;
                                                                                                      DB 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  onse; immunological carrier;
gene therapy; ss.
                                                          8
                                                                                                    Length 1661;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 1573;
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                                                       Indels
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30-APR 1999
04-MAY 1999
05-MAY 1999
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11-MAY 1999
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21-MAY 1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        06-APR-1999;
08-APR-1999;
16-APR-1999;
19-APR-1999;
                                                                                                      10-JUN-1999;
10-JUN-1999;
14-JUN-1999;
  16-JUN-1999;
16-JUN-1999;
17-JUN-1999;
18-JUN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hybridisation assay; genetic mapping; gene expression control; protein identification; signal transduction pathway; metabolic pathway; promoter; termination sequence; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAC47974;
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                                                                                                                                                                                08-JUN-1999
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23-MAR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-FEB-1999;
05-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EP1033405-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Arabidopsis thaliana.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Arabidopsis thaliana DNA fragment SEQ ID NO: 55799
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18-OCT-2000
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99US-0132485

99US-0130479

99US-0130489

99US-0132489

99US-0132485

99US-0132485

99US-01324863

99US-0132487

99US-0132487

99US-0132487

99US-0134287

99US-0134218

99US-0134218

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99US-0134219

99US-0134219

99US-0134219

99US-0135124

99US-0135124

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99US-0135124

99US-0135021

99US-0138045

99US-0138452

99US-0139453

99US-0139453
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99US-0126785
99US-0127462
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99US-0123180.
99US-0123548.
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99US-0128714.
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RESULT 13
AAH56861/c
ID AAH56861
                                                                                             δÃ
                                                                                                                      Query Match 63.6%;
Best Local Similarity 63.6%;
Matches 14; Conservative
                                                                                                                                                                                17-AUG-1999
20-AUG-1999
20-AUG-1999
21-AUG-1999
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24-AUG-1999
27-AUG-1999
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29-SEP-1999
21-CCT-1999
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                                                                 tgttggannnnnnnntttgata
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 standard;
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9908-0149722.
9908-0149922.
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9908-0161105.
 DNA;
  2683
                                                                     976
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                                                                                                                        0;
                                                                                                                        Score 14; DB
Pred. No. 1.4e
0; Mismatches
 ВP
                                                                                                                DB 2-
1.4e+02;
8;
                                                                                                                                                    Length 2660;
                                                                                                                           Indels
                                                                                                                          0;
                                                                                                                          Gaps
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18-JUN 1999
18-JUN

99US-0139456 99US-0139456 99US-0139459 99US-0139461 99US-0139461 99US-0139461 99US-0139462 99US-0139462 99US-01406353 99US-01406353 99US-0140695 99US-014236 99US-014236 99US-014236 99US-014354 99US-014436 99US-0144333 99US-0144335 99US-0144335 99US-0144335 99US-0144335 99US-0144335 99US-0145086 99US-014708 99US-0148386 99US-0148849

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                                                                                                                                                                                               γ
                                                                                                                                                                                                                               cc and antiproliferative activities, and can be used in antisense therapy can defor inhibition of expression of groes or groes. (I) are useful for conhibiting expression of GL or GS in cells or tissues in vitro. (I) are calso useful for inhibiting the growth of a microorganism, or inhibiting conditions of GL or GS gene in a microorganism (a bacterial cell or GS the expression of GL or GS gene which involves administering to the microorganism or to a cell infected with the microorganism, (I). (I) are calso useful for treating a mammalian pathological condition mediated by the microorganisms which involves identifying a cukaryotic organism can be administering (I) such that the growth of microorganism condition mediated by the microorganisms having a GL or GS gene and administering (I) such that the growth of microorganism is inhibited. The antisense compounds are utilised for diagnostics, therapeutics, prophylaxis and as research reagents and kits, e.g., to prevent or delay microbial infections in humans. They are also useful as conceptually microbial infections in humans. They are also useful as conceptually microbial infections in humans and kits, e.g., to prevent or delay microbial infections in humans. They are also useful as completication of the present invention. AAH56855 to AAH56870 represent CC exemplification of the present invention. AAH56855 to AAH56870 represent conditions are used in the present invention.
                                                                                                         Query Match
Best Local Similarity
                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antisense oligonucleotides to nucleotide sequences encoding groE. More generally, antisense compounds (1) comprising antisense oligonucleotides of 5-50 bases targeted to a nucleotide sequence encoding groEL (heat shock protein (HSP)60) (GL) and groES (HSP10) (GS) gene from a microorganism, where the antisense compound is complementary to GL or GS of a microorganism and specifically hybridises with and inhibits the expression of GL or GS, is claimed. (I) have antibacterial, antiviral
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Antisense oligonucleotide; groE; groEL; groES; inhibitor; growth; microorganism; Escherichia coli; Streptcocccus pneumoniae; diagnosis; Streptcocccus pyogenes; Staphylococcus aureus; Pseudomonas aeruginosa; antibacterial; antiviral; antiproliferative; antisense therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Fig 7; 110pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel antisense compounds targeting nucleic acid encoding groEL or groES gene of microorganism, which hybridize with and inhibit expression of the genes, useful to inhibit growth of microorganism
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Streptococcus pyogenes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               microbial infection; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             S. pyogenes groEL gene partial sequence SEQ ID NO:7
                                                                                                                                                                                               Sequence 2683 BP; 808 A; 490 C; 593 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention specifically claims AAH56368 to AAH56832 which are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2001-355633/37.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20-NOV-2000; 2000WO-CA01347.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  having the genes -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (GENE-) GENESENSE TECHNOLOGIES INC
    963
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                         tgttggannnnnnnntttgata
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                                                                                      Conservative
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                                                                                                         63.68;
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                                         22
                                                                                  0;
                                                                                                           Score 14;
Pred. No.
                                                                                    Mismatches
                                                                                                                                                                                             792 T; 0 other;
                                                                                                                                  DB 22;
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                                                                                                                             Length 2683;
                                                                                      Indels
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RESULT 14
AAQ994430/c
ID AAQ994
AX AAQ994
AX AAQ994
AX Surfac
XX Host
FT CDS
FT CDS
FT CDS
FT CDS
FT CDS
FT Sig_pc
FT Mat_pc
FT Mat_pc
FT Host
IN WPI;
DR P-PSDI
XX P-PSDI
XX WPI;
DR P-PSDI
XX WPI;
AAC28622
ID AAC2
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Best Local S
Matches 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A probe based on the N-terminal sequence of B. sphaericus P-1 3855) surface layer protein was used to screen an HindIII-generated library to isolate the slp gene. Promoter regions of the gene are used in genetic constructs providing surface expression of heterologous proteins in P-1 hosts.
                                                                                                                                                                                                                                              2836
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                         06-OCT-2000
                                                                      AAC28622;
                                                                                                                    AAC28622 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 4197 BP; 1470 A; 731 C; 763
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Host cell expressing surface layer protein fusion protein - used for host presentation of antigens and vaccine \operatorname{prod} n.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1995-263827/34.
P-PSDB; AAR80530.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        14-JAN-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        13-JAN-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Bacillus sphaericus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             surface expression; epitope;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22-DEC-1995
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                                                                                                                                                                                                                                           TGTTGGAGCAACTGGTTTGATA
                                                                                                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    layer protein; SLP; fusion protein; vaccine; antigen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Fig.6; 95pp;
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                      (first entry)
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185..3850
/*tag= d
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95..184
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
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                                                                                                                       cDNA; 149
                                                                                                                                                                                                                                                                                                                                                                63.6%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Dhaese
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                DB 16;
1.5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1233 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                         Length 4197;
                                                                                                                                                                                                                                                                                                                                            0;
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Human secreted protein 5'

EST,

SEQ ID NO: 32697.

밁 δÃ Query Match
Best Local Similarity 61.9
Matches 13; Conservative The present sequence is one of a large number of 5' ESTs derived from mRNAs encoding secreted proteins. No ORF has yet been conclusively identified within the present sequence. The 5' ESTs were prepared from total human RNAs or polyA+ RNAs derived from 30 different tissues. EST sequences usually correspond mainly to the 3' untranslated region (UTR) of the mRNA because they are often obtained from oligo-dT primed cDNA libraries. Such ESTs are not well suited for isolating cDNA sequences derived from the 5' ends of mRNAs and even in those cases where longer cDNA sequences have been obtained, the full 5' UTR is rarely included. 5' ESTs are derived from mRNAs with intact 5' ends and can therefore be used to obtain full length cDNAs and genomic DNAs. 5' ESTs are also used in diagnostic, forensic, gene therapy and chromosome mapping procedures. They are used to obtain upstream regulatory sequences and to design expression and socretion vectors. New nucleic acid that is a 5' expressed sequence tag (5' EST) for obtaining cDNAs and genomic DNAs that correspond to 5'ESTs and for diagnostic, forensic, gene therapy and chromosome mapping procedures -Sequence 149 BP; 33 A; 14 C; 37 G; 65 T; 0 other; expression and secretion Claim 1; SEQ ID 32697; 71pp + CD-ROM; English. WPI; 2000-500381/45. Dumas Milne Edwards J, (GEST) GENSET. 26-FEB-1999; 21-FEB-2000; 2000EP-0200610. 06-SEP-2000. EP1033401-A2 Homo sapiens Human; 5' EST; expressed sequence tag; secreted protein; cDNA isolation; gene therapy; chromosome mapping; ss. 13 1 tgttggannnnnnnntttgat 21 tgttggattcttatatttgat 33 99US-0122487 59.1%; Duclert A, vectors. Score 13; DB: Pred. No. 3.4e. 0; Mismatches 0 Giordano J; 3.4e+02; DB 21; Length 149; Indels 0; Gaps 0;

Search completed: December 15, 2001, Job time: 7886 sec

03:29:46

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Sequence

2, Appli 2, Appli 1, Appli 9, Appli 2, Appli 38, Appli 38, Appli 38, Appli 1, Appli 1, Appli

3, Appli 3, Appli 3, Appli 1, Appli 3, Appli 3, Appli 3, Appli 26, Appli

Sequence

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OM nucleic - nucleic search, using sw
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1: /cgn2_6/ptodata/2
2: /cgn2_6/ptodata/2
3: /cgn2_6/ptodata/2
4: /cgn2_6/ptodata/2
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        Minimum Match
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; Sequence 14, Application US/08682517
; Patent No. 5874267
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; Sequence 13, Application

; Patent No. 5874267

; GENERAL INFORMATION:
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; TOPOLOGY: Linear
; MOLECULE TYPE: DNA (ger
US-08-682-517-13
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Best Local S
Matches 14
                                                                                                                                                                                   Patent No. 5874267
GENERAL INFORMATION:
TITLE OF INVENTION: Expression of su:
NUMBER OF SEQUENCES: 25
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, V.
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION:
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 3666 base pairs
TYPE: nucleic acid
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatil
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US-09-037-601-3
PCT-US93-03275-1
PCT-US94-13200-3
US-09-293-322C-9
US-08-121-202-3
US-08-125-587-26
US-08-468-036-2
US-08-981-729-9
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; MOLECULE TYPE:
US-08-682-517-7
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: Sequence 7, Application US/08682517

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Patent No. 5874267
GENERAL INFORMATION:
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SEQUENCE CHARACTERISTICS:
LENGTH: 4197 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 25
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC COMPUTER: PC-DOS/MS-DOS
COERRATING SYSTEM: PC-DOS/MS-DOS
COERRAPEN DATA PD-1006/MS-DOS
TITLE OF INVENTION: Expression of NUMBER OF SEQUENCES: 25
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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STRANDEDNESS: single
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SEQUENCE CHARACTERISTICS:
LENGTH: 4197 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: Patentin Release #1.0, Version #1.30 (EPO) CURRENT APPLICATION DATA:
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APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Word Perfect 6.1 for Windows/MS-DOS 6.2
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/276,531
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2836 TGTTGGAGCAACTGGTTTGATA 2815
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                APPLICANT: Guegler, Karl J.

APPLICANT: Baughn, wariah R.

TITLE OF INVENTION: COMPOSITION FOR THE DETECTION OF GENES ENCODING
TITLE OF INVENTION: RECEPTORS AND PROTEINS ASSOCIATED WITH CELL PROLIFERATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FEATURE:
                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
                                                                         CLASSIFICATION:
PRIOR APPLICATION DATA:
                 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                          STATE: C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: nucleic acid
                                                                                                                                                                                                                                                                          STREET: 3174 POI
CITY: PALO ALTO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INDIVIDUAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRANDEDNESS:
                                CLASSIFICATION:
                                           APPLICATION NUMBER: 60/079, FILING DATE: March 27, 1998
                                                                                                       FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 tgttggannnnnnnntttgata
                                                                                                                                                                                                                                                                                                       ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                14;
                                                                                                                                                                                                                                                                                                                                     OF SEQUENCES:
                                                                                                                                                                                                                               94304
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                                                                                                                                                                                                                                                          CALIFORNIA
Lynn E. Murry, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Application US/09276531
                                                                                                                                                                                                                                                                                        3174 PORTER DRIVE
                                                                                                                                                                                                                                            USA
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                                                                                                                                                                                                                                                                                                                                                                                                                           Yue, Henry
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Hillman, Jennifer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bandman,
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95..3850
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185..3850
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                                                                                                       Herewith
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                                                                                                                                                                                                                                                                                                                                                                                              Karl J
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                                                             60/079,677
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Pred. No.
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33;
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US-08-714-918-19
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; LIBRARY: PROST
; CLONE: 714029
US-09-276-531-32
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Best Local Similarity
                                                                               SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION UNMBER: US/08/714,918
FILING DATE: September 13, 1996
CLASSIFICATION UNMBER: 424
PRIOR APPLICATION DATA:
APPLICATION UNMBER: 60/009,102
FILING DATE: December 22, 1995
APPLICATION NUMBER: 60/003,798
APPLICATION NUMBER: 60/003,798
FILING DATE: September 15, 1995
APPLICATION UNMBER: 50/003,798
FILING DATE: September 15, 1995
APPLICATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 32,327
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: (650) 845-4166
INFORMATION FOR SEQ ID NO: :
SEQUENCE CHARACTERISTICS:
LENGTH: 3487 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
NFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" biskette, 1.44 Mb
MEDIUM TYPE: storage
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REFERENCE/DOCKET NUMBER: PA
TELECOMMUNICATION INFORMATION:
                                                 TELECOMMUNICATION INFORMATION: TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: STAPHYLOCOCCUS AUREUS ANTIBACTERIAL ITTLE OF INVENTION: TARGET GENES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM CONTROL OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: Los Angeles
STATE: California
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                               TELEFAX:
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STRANDEDNESS: sing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE:
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633 West Fifth Street
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Martin, Patrick K.
Schmid, Molly B.
                                 (213)
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                                                                                                                                                                                                                                                                                                                                  IBM Compatible
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                                                                                                                                                                                                                                                  UMBER: US/08/714,918
September 13, 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Dongxu
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                              955-0440
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                                                                                   222/005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 13;
Pred. No. 1
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; LENGTH: 5253 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
US-08-714-918-19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 19, Application US/09265315 Patent No. 6187541
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                                                         INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                   FILING DATE: December 22, 19
APPLICATION NUMBER: 60/003,7
FILING DATE: September 15, 1
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/
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                                                                                                                                        REFERENCE/DOCKET NUMBER: 24
TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
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STREET: Suite 4700
CITY: Los Angeles
STATE: California
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                                                                                                                                                                                                                                                               APPLICATION NUMBER: 60/
                                                                                          TELEFAX: (213)
TELEX: 67-3510
                                                                                                                                                                                                                                                                                                                               CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                              FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                         OPERATING SYSTEM:
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                STRANDEDNESS:
                                                                                                                       TELEPHONE:
                                                                                                                                                                                                                                                                                             APPLICATION NUMBER:
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                                             LENGTH:
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                        nucleic acid
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Lee, Ving J.
'-louin, Francois
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                                                                                                                                                                                                                                                                                                                                                                                                                           IBM Compatible
linear
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                                                                                                                          (213) 489-1600
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March 9, 1999
เพ. ภาค
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                                                                                                          955-0440
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61.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             METHODS OF SCREENING FOR COMPOUNDS ACTIVE ON STAPHYLOCOCCUS AUREUS
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mber 13, 1996
60/009,102
                                                                                                                                                                                                                                  60/003,
                                                                                                                                                              32,327
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Pred. No. 1.2e+02;
                                                                                                                                                        240/247
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                                       Query Match
Best Local Similarity
Watches 13; Conserv
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                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM Compatible
ODERATING SYSTEM; IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/265,315
FILING DATE: March 9, 1999
CLASSIFICATION LOATA:
APPLICATION UNMBER: 08/714,918
FILING DATE: September 13, 1996
APPLICATION NUMBER: 60/009,102
FILING DATE: December 22, 1995
APPLICATION NUMBER: 60/003,798
FILING DATE: September 15, 1995
APPLICATION NUMBER: 60/003,798
FILING DATE: September 15, 1995
APPLICATION NUMBER: 60/003,798
FILING DATE: MERCHARTION:
NAME: WAIDLING, Richard J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: Suite
STREET: Suite
STREET: LOS Angeles
California
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APPLICANT:
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MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2895 GTTGGATGTAGTCTTTTGATA 2915
2895 GTTGGATGTAGTCTTTTGATA 2915
                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION:
TITLE OF INVENTION:
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nes 13; Conserv
                                                                                                                                                            STRANDEDNESS:
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                                                                                                                                                                                          LENGTH:
               gttggannnnnnnntttgata 22
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o. 6187541
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67-3510
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633 West Fifth Street
Suite 4700
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Martin, Patrick K.
Schmid, Molly B.
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Malouin, Francois
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                                                      Conservative
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                                                                                                                                                linear
                                                                                                                                                            single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 METHODS OF SCREENING FOR COMPOUNDS ACTIVE ON STAPHYLOCOCCUS AUREUS TARGET GENES
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61.9%;
                                                                    59.1%;
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                                                                                                                                                                                                                                                                                                           32,327
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Pred. No. 1.2e+02;
                                                        0;
                                                                                                                                                                                                                                                                                                 240/247
                                                                      Score 13;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1.44 Mb
                                                        Mismatches
                                                  DB 4; Lc.., 1.2e+02; 8;
                                                                                   Length 5253;
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                                                        Indels
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RESULT 9
US-09-266-417-19
RESULT 10
US-08-821-994-61/c
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                                                                                                                                      Best
                                                                                                                        Matches
                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                          INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: September 15, 1995 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage
                                                                                                                                                                                                                                                                                                                                                     NAME: Watburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 24(
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
                                                            2895 GTTGGATGTAGTCTTTTGATA 2915
                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER:
FILING DATE: September:
APPLICATION NUMBER:
FILING DATE: Decembe
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
                                                                                                                                                                                                        TYPE: ...
STRANDEDNESS: SING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM Compatible OPERATING SYSTEM: IBM P.C. SOFTWARE: Word Perfect 5.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US FILING DATE: March 9,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET:
                                                                                                                       Local Similarity 61.9 les 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: U.S.A. ZIP: 90071-2066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE:
                                                                                                                                                                                                                                                              LENGTH:
                                                                                                                                                                                                                                                                                                                        TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
                                                                            2 gttggannnnnnnntttgata 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION:
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California
                                                                                                                                                                                                                                               nucleic acid
                                                                                                                                                                                                                                                                                                            67-3510
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633 West Fifth Street
Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Application US/09266417
                                                                                                                                                                                                                                                          5253 base pairs
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Malouin, Francois
Martin, Patrick K.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Benton, Bret
                                                                                                                                                                                                                                                                                                                                                                                                                                                            September 13, 1990
NUMBER: 60/009,102
December 22, 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           METHODS OF SCREENING FOR ACTIVE ON STAPHYLOCOCCUS
                                                                                                                                     59.1%;
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), 1999
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                                                                                                                        0;
                                                                                                                                   Score 13; DB 4;
Pred. No. 1.2e+02;
                                                                                                                        Mismatches
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                                                                                                                                                     Length 5253;
                                                                                                                        0;
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Sequence 61, Application US/08821994A Patent No. 6228643

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GENERAL INFORMATION:

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US-08-413-118-118
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; ORGANISM: Brassica napus
US-08-821-994-62
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; ORGANISM: Brassica napus
US-08-821-994-61
                                                                                                                                                           RESULT 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: Promoters
FILE REFERENCE: PPD 50108
CURRENT APPLICATION NUMBER: US/08/821,994A
CURRENT FILING DATE: 1997-03-22
EARLIER APPLICATION NUMBER: PCT/GB97/00729
EARLIER FILING DATE: 1997-03-18
EARLIER FILING DATE: 1997-03-18
EARLIER FILING DATE: 1996-03-22
NUMBER OF SEQ ID NOS: 89
CONTENDED: DATE: 1996-03-22
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FILE REFERENCE: PPD 50108

CURRENT APPLICATION NUMBER: US/08/821,994A

CURRENT FILING DATE: 1997-03-22

EARLIER APPLICATION NUMBER: PCT/GB97/00729

EARLIER FILING DATE: 1997-03-18

EARLIER APPLICATION NUMBER: GB 9606062.9

EARLIER FILING DATE: 1996-03-22

NUMBER OF SEQ ID NOS: 89
                                                                                Sequence 118, Application US/08413118 Patent No. 5688920 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQ ID NO 62
LENGTH: 1434
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Sequence 62, App...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
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Best Local
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LENGTH: 1390
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Greenland, Andrew J. APPLICANT: Thomas, Didier RP APPLICANT: Jepson, Ian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Greenland, Andrew J
APPLICANT: Thomas, Didier RP
APPLICANT: Jepson, Ian
                                              APPLICANT: PAOLETTI, APPLICANT: LIMBACH,
                TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                       Local Similarity nes 13; Conserv
                                                                                                                                                                                                                26 TGTTGGTTTTCTAGTTTTGATA 5
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                                                                                                                                                                                                                                                                                       Conservative
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                                                                 PAOLETTI, ENZO
                                                                                                                                                                                                                                                                                                     56.4%;
59.1%;
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         KEITH J.

NUCLEOTIDE AND AMINO ACID SEQUENCES OF
CANINE HERPESVIRUS 9B, 9C, AND 9D AND USES THEREFOR
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Pred. No. 2
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Pred. No. 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
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; TOPOLOGY: linear; MOLECULE TYPE: cDNA
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                                                                                                                                                                                                                                                                                                                                                                                                                            Patent No. 601754
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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TELEPHONE: (212) 840-3333
                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 30-MAR-1994
ATTORNEY/AGENT INFORMATION:
NAME: FROMMER, WILLIAM S.
REGISTRATION NUMBER: 25,506
REFERENCE/DOCKET NUMBER: 454310-2670
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0
FILING DATE: 30-MAR-1994
              CLASSIFICATION: PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: PAOLETTI, ENZO APPLICANT: LIMBACH, KEITH J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION:
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                                                    FILING DATE
                                                                                                                                                                                                               COUNTRY:
                                                                                                                                                                                                                                                    CITY: NEW YORK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: CLASSIFICATION:
                                                                   APPLICATION NUMBER:
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APPLICATION NUMBER:
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: UNITED STATES OF AMERICA
                                                                                                                                                                                                                                 NEW YORK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1760 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             E: CURTIS, MORRIS & SAFFORD, 530 FIFTH AVENUE, 25TH FLOOR
                                                                                                                                                                                                                                                                   E: CURTIS, MORRIS & SAFFORD, P.C
530 FIFTH AVENUE, 25TH FLOOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                               UNITED STATES OF AMERICA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       linear
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59.1%;
                                                                                                                                                                                                                                                                                                                                      NUCLEOTIDE AND AMINO ACID SEQUENCES CANINE HERPESVIRUS gB, gC, AND gD AN
                                                                                                                                                                                                                                                                                                                         128
08/413,118
                                                                     US/08/473,446
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Pred. No. 2.2e+02
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 1;
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                                                                                                                                                                                                                                                                                                                                        gD AND USES THEREFOR
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Best Local Similarity 59.1
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 2, Application US/08752238 Patent No. 5804418
                                                                                                                                            TELEFAX: (216) 241-0816 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (212) 840-0712 INFORMATION FOR SEQ ID NO: 118:
                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1076 TGTTGGATAACTATTTTTATA 1097
                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                 FEATURE:
                                                                                                                             SEQUENCE CHARACTERISTICS:
                                                                                                                                                                          TELECOMMUNICATION INFORMATION: TELEPHONE: (216) 622-8458
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 12
CORRESPONDENCE ADDRESS:
ADDRESSEE: CALFEE, HALTER & GRISWOLD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT:
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                                               MOLECULE TYPE:
                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
NAME: GOLrick, Mary E.
REGISTRATION NUMBER: 34,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE: CDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION:
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ATTORNEY/AGENT INFORMATION:
NAME/KEY:
                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: 246
ILECOMMUNICATION -
                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: 800 Supr
CITY: Cleveland
STATE: Ohio
                                                                                     TYPE: nucleic acid
                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/08/752,238 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 1760 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                  TOPOLOGY:
                                                                            STRANDEDNESS:
                                                                                                                                                                                                                                                                   CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: UZIP: 44114
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REFERENCE/DOCKET NUMBER: 45
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             800 Superior Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matsuura Dr., Manabu
WENTION: Method for Preparing Nucleotide
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Mohr Dr., Georg
Saldanha Dr., Roland
                                                                  linear
                                                                                                                                                                                                                                                                                                                                                                              Floppy disk
                                              DNA (genomic)
                                                                            double
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59.18;
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Pred. No. 2
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2.2e+02;
9;
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                                                          Matches
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                                                                     Query Match
Best Local Similarity
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Best Local
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TELEPHONE: (216) 622-8416
TELEPAX: (216) 241-0816
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Beall Dr., Clifford J.
TITLE OF INVENTION: Methods of Making an Rnp Particle
TITLE OF INVENTION: Having Nucleotide Integrase Activity
                                                                                                                                                                                                                                                                                                                                                                              FILING DATE:
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                          FEATURE:
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139 TATTGGAATATAAATTTTGATA 118
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CITY: Cleveland
                                                                                                                                                              NAME/KEY:
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                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER:
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                       1 tgttggannnnnnnntttgata 22
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Mohr Dr., Georg
Saldanha Dr., Roland
                                                       Conservative
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Zimmerly Dr., Steven
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                                                                                                                                                                                        DNA (genomic)
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                                                                     56.4%;
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                                                                     Score 12.4;
Pred. No. 2.
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                                                        Mismatches
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                                                    2.2e+02;
9;
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                                                       Gaps
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Search completed: December 15, 2001, 03:22:49
Job time: 7564 sec

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq
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Perfect score:
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length: 2000000000
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Gapop 10.0 , Gapext 1.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         11351937 seqs, 5372889281 residues
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22
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em_htc:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	score	Query Match Length DB	ength	DB	ID	Description
c 1	14	63.6	184	10	вв357781	вв357781
2	14	63.6	210	10	BE059640	BE059640
c 3	14	63.6	228	11	BE956026	BE956026
c 4	14	63.6	235	10	BB170254	вв170254
ი ა	14	63.6	235	13	AZ925147	AZ925147
6	14	63.6	250	10	AI122094	AI122094
c 7	14	63.6	261	10	AW879900	AW879900 QV3-OT002
c 8	14	63.6	265	10	BB287901	вв287901
ი 9	14	63.6	269	H	BF800082	BF800082
c 10	14	63.6	293	11	BF634128	BF634128
11	14	63.6	298	10	AA515145	AA515145
12	14	63.6	332	10	ロロンのこうちん	BE303756

Email: genome-res@gsc.riken.go.jp,

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498	497	496	493	490	489	485	482	481	476	471	467	459	447	436	433	427	416	416	410	410	399	389	386	378	376	376	371	364	359	349	342	336
11	10	13	10	10	10	11	13	11	13	10	10	11	13	11	11	10	13	10	13	10	11	10	11	10	13	13	13	10	10	10	13	13
F00332	BE353412	8982	AA637362	BE448206	9682	BF226581	в29252	T75954	1460	AI591928	AL370583	BE854379	AZ273408	BF463084	BF464046	AI662442	AQ549780	AA555292	AQ150667	158	R89220	AA488558	BF318641	AW240466	AQ102282	AQ844476	AQ311167	AW226052	AV546877	₽	BH083614	AZ037213
BF003327 EST431825	BE353412 EST353789	AQ589820 HS_2136_B	AA637362 vu09f03.r	BE448206 ut80g10.y	BE196823 ug71a09.y	BF226581 uz12f09.x	B29252 T29J13TF TA		nbeb0049	mt31b03	MtBA38	ux29e12	8 RPCI-23-	4 UI-M-C	4046 UI-M-CG0	<pre>mt25a11.</pre>) RPCI-11	nk82g0	' HS_3197_	EST322	p99a12.r	8 ab37h10.	ux29e1	٠.	PHS_302	5 an37g1		ST76G0	' AV54687	8 AU1129	614 RPCI-24-	AZ037213 RPCI-23-3

ALIGNMENTS

) T I I I V C	TITLE R JOURNAL U COMMENT C	· · · · · · × × × × × × × × × × × × × ×	REFERENCE 1 AUTHORS K	ACCESSION B VERSION B KEYWORDS E SOURCE h	RESULT 1 BB357781/c LOCUS DEFINITION B
LADOTATORY FOR GENOME EXPLORATION RESEATOR GETOUP, KIKEN GENOMIC Sciences Center(GSC), Yokohama Institute of Physical and Chemical Research (RIKEN) 1-7-22 Suehiro-cho, Tsurumi-ku, Yokohama, Kanagawa 230-0045, Japan Tel: 81-45-503-9212 Fax: 81-45-503-9216	,M., Muchmatsu,M. and Hayashizaki,Y. RIKEN MOUSE EST'S (Konno,H., et al.) Unpublished (2000) Contact: Yoshhide Hayashizaki	Izawa, M., Kadota, K., Kagawa, I., Kai, C., Kawai, J., Kikuchi, N., Izawa, M., Kadota, K., Kagawa, I., Kai, C., Kawai, J., Kikuchi, N., Kiyosawa, H., Kojima, Y., Kondo, S., Koya, S., Kurihara, C., Kusakabe, M., Matsuyama, T., Miki, R., Mizuno, Y., Nakamura, M., Oda, H., Okazaki, Y., Ono, T., Owa, C., Saito, H., Sakai, C., Sato, K., Shibata, K., Shibata Y., Shigemoto, Y., Shinagawa, A., Shiraki, T., Sogabe, Y., Sugahara, Y., Suzuki, H., Suzuki, H., Tagawa, A., Takahashi, F., Tominaga, N., Toya T., Tsunoda, Y., Watahiki, A., Watanabe, S., Yamamra, T., Yamanaka, I. Yano, R., Yasunishi, A., Yokota, T., Yoshida, K., Yoshiki, A., Yoshino	Ebkaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (bases 1 to 184) Konno, H., Aizawa, K., Akahira, S., Akiyama, J., Arakawa, T., Carninci Konno, T., Fukuda, S., Fukunishi, Y., Hara, A., Hayatsu, N., Tith Mara, M., M	BB357781.1 GI:9069609 EST. house mouse. Mus musculus	

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BASE COUNT
ORIGIN
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VERSION
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                                                             SOURCE
ORGANISM
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Carninci,P., Nishiyama,Y., Westover,A., Itoh,M., Nagaoka,S., Sasaki
N., Okazaki,Y., Muramatsu,M. and Hayashizaki,Y.
Phermostabilization and thermoactivation of thermolabile enzymes by trehalose and its application for the synthesis of full length cDNA. Proc. Natl. Acad. Sci. U.S.A. 95 (2), 520-524 (1998)
Itoh,M., Kitsunai,T., Akiyama,J., Shibata,K., Izawa,M., Kawai,J., Tomaru,Y., Carninci,P., Shibata,Y., Ozawa,Y., Muramatsu,M., Okazaki
Y and uwashi zai v
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Carninci,P. and Hayashizaki,Y.
High-efficiency full-length cDNA cloning. Methods Enzymol. 303,
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                                                                                                                                                                                        BE059640 210 bp mRNA EST 21-NOV-2000 sn35a01.yl Gm-c1016 Glycine max cDNA clone GENOME SYSTEMS CLONE ID: Gm-c1016-13441 5' similar to TR:Q9ZVI3 Q9ZVI3 PUTATIVE ELONGATION
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosidae; eurosids I; Fabales; Fabaceae; Papilionoideae; Phaseoleae
                                                             Glycine max
                                                                                                                                                 BE059640
                                                                                                                                                                      FACTOR. ;, mRNA sequence.
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14; Conservative
                                                                                                                               BE059640.1 GI:8404006 ·
                                                                                      soybean
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              visit our web site (http://genome.rtc.riken.go.jp) for
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/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="C030024G02"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               prepared and sequenced in Mouse Genome Encyclopedia prepared of Genome Exploration Research Group in Riken Project of Genome Exploration Research Group in Riken Genomic Sciences Center and Genome Science Laboratory in Genomic Sciences Center and Genome Science Laboratory in RIKEN Division of Experimental Animal Research in Riken RIKEN Division of Experimental Animal Research in Riken
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /clone_lib="RIKEN full-length enriched, adult male corpus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /tissue_type="corpus striatum"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /sex="male"
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Washington University School of Medicine
4444 Forest Park Parkway, Box 8501, St. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tel: 314 286 1800
Fax: 314 286 1810
Email: est@watson.wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Contact: Shoemaker R/Public Soybean
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             info@genomesystems.com web site: www.genomesystems.com
Insert Length: 647 Std Error: 0.00
Seq.primer: -40RE from Gibco.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This clone is available through: Genome Systems, Inc. 4633 World Parkway Circle St. Louis, Missouri 63134 For further information call: (800) 430-0030 or (314) 427-3222 FAX:(888) 919-3324 or (314) 427-3324 or contact: clones@genomesystems.com or
                                                                                                                                                                            BE956026 228 bp mRNA EST 04-OCT-2000 UI-M-BH4-bav-a-11-0-UI.S1 NIH_BMAP_M_S5 Mus musculus cDNA clone UI-M-BH4-bav-a-11-0-UI 3', mRNA sequence.
Mus muscullus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Eukaryota; Metazoa; Rodentia; Sciurognathi; Muridae; Murinae; Mus
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
1 (bases 1 to 228)
                                                                                                                                    BE956026.1
                                                                                                                                                         BE956026
                                                                                          house mouse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note="Vector: pBluescript II XR: Site_1: EcoRI; Site_2: /note="Vector: pBluescript II XR: Site_1: EcoRI; Site_2: XhoI; This cDNA library was constructed from mRNA isolated from immature flowers of field grown plants. The cDNA library was prepared using the Stratagene pBluescript II XR library construction kit. Complementary DNA was XR library construction kit. Complementary DNA was ynthesized from mRNA using a primer consisting of a poly (dT) sequence with a XhoI restriction site. EcoRI adapters were ligated to the blunt-ended cDNA fragments followed by XhoI digestion. The cDNA fragments were directionally cloned into the EcoRI-XhoI restriction site of the pBluescript vector. The ligated cDNA fragments were transformed into XLIO-Gold host cells. This library was constructed by Dr. Randy Shoemaker and Dr. John Erpelding."

57 a 38 C 55 g 60 t
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/db_xref="taxon:3847"
/clone="GENOME SYSTEMS CLONE ID: Gm-c1016-13441"
/clone_lib="Gm-c1016"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /tissue_type="immature flowers of field grown plants"
/lab_host="XL10-Gold"
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Pred. No. 2.1e+03;
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RESULT 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 tgttggannnnnnntttgata 22
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National Institute of Mental Health
                                                                                                                            BB170254 235 bp mRNA EST 29-JUN-2000 BB170254 RIKEN full-length enriched, adult male hypothalamus Mus musculus cDNA clone A230017D13 3', mRNA sequence.
BB170254 BB170254 1 GI:8829337 EST.
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Fax: 301 443 9890
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                                            Mus musculus
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                             house mouse.
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/Clone_1ib="NIH_BMAP_M_SS"
/Clone_1ib="NIH_BMAP_M_SS"
/dev_stage="27,32 days"
/dev_stage="27,32 days"
/lab_host="DH10B (Life Technologies)"
/lab_host="DH10B 
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TAG_SEQ-CAAGG":
47 c 42 g 75
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TAG_LIB=NIH_BMAP_M_S5
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/strain="C57BL/6J"
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Respondent to 235, Fukunishi,Y., Hara,A., Hayatsu,N.,

P., Endo,T., Fukuda,S., Fukunishi,Y., Hara,A., Hayatsu,N.,

Hirozane,T., Hori,F., Ishii,Y., Ishikawa,J., Ishikawa,T., Itoh,M.,

Izawa,M., Kadota,K., Kagawa,I., Kai,C., Kawai,J., Kikuchi,N.,

Kiyosawa,H., Kojima,Y., Kondo,S., Koya,S., Kurihara,C., Kusakabe,M.,

Mitsuyama,T., Miki,R., Mizuno,Y., Nakamura,M., Oda,H., Okazaki,Y.,

Ono,T., Owa,C., Saito,H., Sakai,C., Sato,K., Shibata,K., Shibata,Y.,

Sundi,H., Saito,H., Sakai,C., Sato,K., Shibata,Y.,

Sundi,H., Suzuki,H., Sakai,C., Sato,K., Shibata,Y.,

Suzuki,H., Suzuki,H., Tagawa,A., Shiraki,F., Tominaga,N., Toya,

T., Tsunoda,Y., Watahiki,A., Watanabe,S., Yamamura,T., Yamanaka,I.,

Yano,R., Yasunishi,A., Yokota,T., Yoshida,K., Yoshiki,A., Yoshino,

M.K., Muzamatsu,M. and Hayashizaki,Y.

RIKEN, Mouse ESTs (Konno,H., et al.)

Unpublished (2000)

Contact: Yoshinde Hayashizaki, T., Patanaba, T., Taganaka, T., Taganaba, T., Taganaba, T., Yamanaka, T., Yamanaka, T., Yamanata, Y., Yamanata, Y., Yanganata, Y.
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URL:http://genome.gsc.riken.go.jp/
Carninci,P., Nishiyama,Y., Westover,A., Itoh,M., Nagaoka,S., Sasaki
,N., Okazaki,Y., Muramatsu,M. and Hayashizaki,Y.
Thermostabilization and thermoactivation of thermolabile enzymes by
Trehalose and its application for the synthesis of full length
cDNA. Proc. Natl. Acad. Sci. U.S.A. 95 (2), 520-524 (1998)
Itoh,M., Kitsunai,T., Akiyama,J., Shibata,K., Izawa,M., Kawai,J.,
Tomaru,Y., Carninci,P., Shibata,Y., Ozawa,Y., Muramatsu,M., Okazaki
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Automated filtration based high-throughput plasmid preparation system. Genome Res. 9 (5), 463-470 (1999)
Carninci,P. and Hayashizaki,Y.
High-efficiency full-length cDNA cloning. Methods Enzymol. 303,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Laboratory for Genome Exploration Research Group, RIKEN Genomic Sciences Center(GSC), Yokohama Institute
The Institute of Physical and Chemical Research (RIKEN)
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Fax: 81-45-503-9216
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/dev_stage="adult"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /clone_lib="RIKEN full-length enriched, adult male
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /clone="A230017D13"
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                                                                                                                       CDNA was cleaved with XhoI and BamHI. Vector: a fied pBluescript KS(+) after bulk excision from Lambda
                                      31 g
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Query Match

63.6%;

Score 14;

DB 10;

Length 235;

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AZ925147/c
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AI122094
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ORGANISM
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Best Local
                                             AUTHORS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ISM Saccharomyces paradoxus
Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
Saccharomycetales; Saccharomycetaceae; Saccharomyces.

CE 1 (bases 1 to 235)
Cliften, P.F., Hillier, L.W., Fulton, L., Graves, T., Miner, T., Gish, W.R., Waterston, R.H. and Johnston, M.
Surveying Saccharomyces genomes to identify functional elements by comparative DNA sequence analysis
AL Unpublished (2001)
Contact: Johnston M
Department of Genetics
Washington University Medical School
Box 8332, 4566 Scott Ave., St. Louis, MO 63110, USA
Tel: 314 362 7855
Fax: 314 362 7855
Email: mj@genetics:wustl.edu
Email: mj@genetics:wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TGTTGGACTTAGTTCTTTGATA 45
                                                                                                                                                                                                                                                                                                                              tgttggannnnnnnntttgata
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4910.ez32e09.s1 Saccharomyces paradoxus N17 genomic clone 4910.ez32e09.s1, DNA sequence AZ925147
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AZ925147.1 GI:13496046
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Class: random plasmid subclone.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                      14; Conservative
                                                                                                                                                                                                  AII22094 250 bp mRNA EST 02-SEP-1998 uc46f10.rl Soares_mammary_gland_NMLMG Mus musculus CDNA clone IMAGE:1401067 5' similar to SW:Y335_MYCGE p47577 HYPOTHETICAL
1 (bases 1 to 250)
Marra, M., Hillier, L., Allen, M., Bowles, M., Dietrich, N., Dubu, Geisel, S., Kucaba, T., Lacy, M., Le, M., Martin, J., Morris, M., Schellenberg, K., Steptoe, M., Tan, F., Underwood, K., Moore, B., Theising, B., Wylie, T., Lennon, G., Soares, B., Wilson, R. and
                                                                                                                                         EST
                                                                                                                                                                      GTP-BINDING PROTEIN MG335. ;, mRNA sequence. ai122094
                                                                            Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                          Mus musculus
                                                                                                                          house mouse
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             /clone="4910.ez32e09.s1"
/clone_lib="Saccharomyces paradoxus N17"
/note="Random genomic sequence"
35 c 36 g 85 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /organism="Saccharomyces
/strain="N17"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /db_xref="taxon:27291"
                                                                                                                                                         GI:3522418
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0; Mismatches
                                                                            Chordata; Craniata; Vertebrata; Euteleostomi;
Rodentia; Sciurognathi; Muridae; Murinae; Mus
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                                                                                                                                                                                                                                                                                                                                            22
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Pred. No.
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ches 8;
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ches 8;
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Best Local
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Contact: Marra M/Mouse EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Waterston,R.
The WashU-HHMI Mouse EST Project
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Email: mouseest@watson.wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                           EST
                                                                                                                                                                      Simpson, A.J.
                                                                                                                                                                                                                                                                                                                                                        AW879900.1
                                                                                                                                       sequence tags
                                                                                         Contact: Simpson A.J.G.
                                                                                                                                                                                                                                                                                                                              nman
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                        Natl. Acad. Sci. U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /organism="Mus musculus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /db_xref="taxon:1
                                                                                                                                                                                                                                                                                                                                                         GI:8041910
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Pred. No.
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Washu HHMI Mouse EST Project
Washington University School of MedicineP
4444 Forest Park Parkway, Box 8501, St. Lo
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High quality sequence stop: 225.
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AW879900 261 bp mRNA EST 23-MAY-2000
QV3-OT0029-290300-134-b02 OT0029 Homo sapiens cDNA, mRNA sequence.
AW879900
                                                                                                                                                                                                                                                                                                                         1 (bases 1 to 261)
Dias Neto,E., Garcia Correa,R., Verjovski-Almeida,S., Briones,M.R.,
Dias Neto,E., Garcia Correa,R., Bordin,S., Costa,F.F.,
Nagai,M.A., da Silva,W. Jr., Zago,M.A., Bordin,S., Costa,F.F.,
Goldman,G.H., Carvalho,A.F., Matsukuma,A., Baia,G.S., Simpson,D.H.,
Goldman,G.H., Carvalho,A.F., Bucher,P., Jongeneel,C.V., O'Hare
Brunstein,A., deOliveira,P.S., Bucher,P., Jongeneel,C.V., O'Hare
M.J., Soares,F., Brentani,R.R., Reis,L.F., de Souza,S.J. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                              Shotgun sequencing of the human transcriptome with ORF
Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
Rua Prof. Antonio Prudente 109, 4 and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note="Vector: pT7T3D-Pac (Pharmacia) with a modified /note="vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; 1st strand cDNA was prepared from mammary polylinker; 1st strand cDNA was prepared from mammary polylinker; 1st stranded cDNA was with a Not I - oligo(dT) primer. Double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT7T3 vector. Library is normalized. Library was constructed by Bento Soares and M. Fatima Bonaldo. " a 79 c 75 g 48 t
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/clone_lib="Soares_mammary_gland_NMLMG"
/sex="female (lactating)"
/tissue_type="mammary gland"
/lab_host="DH10B"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                 97 (7),
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2.1e+03;
              4 andar, 01509-010, Sao Paulo-SP
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                                                                                                                                                                                                                                                                                                                                        Konno, H., Aizawa, K., Akahira, S., Akiyama, J., Arakawa, T., Carninci P., Endo, T., Fukuda, S., Fukunishi, Y., Hara, A., Hayatsu, N., P., Endo, T., Fukuda, S., Fukunishi, Y., Ishikawa, J., Ishikawa, T., Itoh, M., Izawa, M., Kadota, K., Kagawa, I., Kai, C., Kawai, J., Kikuchi, N., Kiyosawa, H., Kojima, Y., Kondo, S., Koya, S., Kurihara, C., Kusakabe, M., Matsuyama, T., Miki, R., Mizuno, Y., Nakamura, M., Oda, H., Okazaki, Y., Ono, T., Owa, C., Saito, H., Sakai, C., Sato, K., Shibata, K., Shibata, Y., Shigemoto, Y., Shinagawa, A., Shiraki, T., Sogabe, Y., Sugahara, Y., Suzuki, H., Suzuki, H., Tagawa, A., Takahashi, F., Tominaga, N., Toya, T., Tsunoda, Y., Watahiki, A., Watanabe, S., Yamamura, T., Yamanaka, I., Yano, R., Yasunishi, A., Yokota, T., Yoshida, K., Yoshiki, A., Yoshino, M., Muramatsu, M. and Hayashizaki, Y., RIKEN Mouse ESTS (Konno, H., et al.)
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Email: genome-res@gsc.riken.go.jp,
URL:http://genome.gsc.riken.go.jp/
Carninci,P., Nishlyama,r., Westover,A., Itoh,M., Nagaoka,S., Sasaki
,N., Okazaki,Y., Muramatsu,M. and Hayashizaki,Y.
Thermostabilization and thermoactivation of thermolabile enzymes by
                                                                                                                                                                             Laboratory for Genome Exploration Research Group, RIKEN Sciences Center(GSC), Yokohama Institute
The Institute of Physical and Chemical Research (RIKEN)
1-7-22 Suehiro-cho, Tsurumi-ku, Yokohama, Kanagawa 230-Tel: 81-45-503-9222
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BB287901 265 bp mRNA
BB287901 RTKEN full-length enriched,
clone B020022N08 3', mRNA sequence.
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                                                                                                                                                        Fax: 81-45-503-9216
                                                                                                                                                                                                                                                                                                                    Contact: Yoshihide Hayashizaki
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; 
Mammalia; Eutheria;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note-"Organ: ovary; Vector: puc18; Site_1: SmaI; Site_2: SmaI; A mini-library was made by cloning products derived from ORESTES PCR (U.S. Letters Patent application No. 196,716 - Ludwig: Institute for Cancer Research) profiles into the pUC 18 vector: Reverse transcription of tissue mRNA and cDNA amplification were performed under low stringency conditions."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /dev_stage="Adult"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Chordata; Craniata; Vertebrata; Euteleostomi; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22
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Pred. No.
                                                                                                                                                                                                                                                                                      Exploration Research Group, RIKEN Genomic
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                                                                                                                         1 (bases 1 to 269)
Dias Neto,E., Garcia Correa,R., Verjovski-Almeida,S., Briones,M.R.,
Nagai,M.A., da Silva,W. Jr., Zago,M.A., Bordin,S., Costa,F.F.,
Goldman,G.H., Carvalho,A.F., Matsukuma,A., Baia,G.S., Simpson,D.H.,
Brunstein,A., deOliveira,P.S., Bucher,P., Jongeneel,C.V., O'Hare
,M.J., Soares,F., Brentani,R.R., Reis,L.F., de Souza,S.J. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Automated filtration-based high-throughput plasmid preparation system. Genome Res. 9 (5), 463-470 (1999)
Carninci,P. and Hayashizaki,Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  trehalose and its application for the synthesis of full length cDNA. Proc. Natl. Acad. Sci. U.S.A. 95 (2), 520-524 (1998)
Itoh, M., Kitsunai, T., Akiyama, J., Shibata, K., Izawa, M., Kawai, J., Tomaru, Y., Carninci, P., Shibata, Y., Ozawa, Y., Muramatsu, M., Okazaki, Y. and Hayashizaki, Y.
                                                                                                                                                                                                                                                                                                                                                                                                             BF800082 269 bp mRNA EST 12-JAN-2001 MR1-CI0021-161000-004-f02 CI0021 Homo sapiens cDNA, mRNA sequence.
Contact: Simpson A.J.G.
                                                              sequence tags
                                                                                Shotgun sequencing of the human transcriptome with ORF expressed
                                                                                                         Simpson, A.J
                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               High-efficiency full-length cDNA cloning. Methods Enzymol. 303,
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/note="Site_1: SalI; Site_2: BamHI; cDNA library was note="Site_1: SalI; Site_2: BamHI; cDNA library was prepared and sequenced in Mouse Genome Encyclopedia Project of Genome Exploration Research Group in Riken Genomic Sciences Center and Genome Science Laboratory in RIKEN. Division of Experimental Animal Research in Riken
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /clone_lib="RIKEN full-length enriched,
/tissue_type="egg"
/dev_stage="2 cells"
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                                           Acad. Sci. U.S.A.
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This sequence was derived from the FAPESP/LICR Human Cancer Genome
Project. This entry can be seen in the following URL
(http://www.ludwig.org.br/scripts/gethtml2.pl?tl=MR1&t2=MR1-CI0021-
161000-004-f02&t3=2000-10-16&t4=1)
Seq.primer: puc 18 forward
Seq.primer: puc 18 forward
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High quality sequence stop: 269.
Location/Qualifiers
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Rua Prof. Antonio Prudente 109, 4 andar, 01509-010, Sao Paulo-SP,
                                                                                                                                                                                                                                                                                                       Torrez-Jerez,I., Scott,A.D., Harris,A.R., Gonzales,R.A., Bell,C.J. Flores,H.R., Inman,J.T., Weller,J.W. and May,G.D. Expressed Sequence Tags from the Samuel Roberts Noble Foundation Medicago truncatula drought library Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 5', mRNA sequence.
BF634128
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NF082F04DT1F1042 Drought Medicago truncatula cDNA clone NF082F04DT
                                                                                                                                                                          Plant Biology Division
The Samuel Roberts Noble Foundation
2510 Sam Noble Parkway, Ardmore, OK 73402, USA
Tel: 580 221 7391
Fax: 580 221 7380
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Medicago truncatula
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
Rosidae; eurosids I; Fabales; Fabaceae; Papilionoideae; Trifolieae
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14; Conservative
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                                                                                                                             Email: gdmay@noble.org
Insert Length: 293 S
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                                                                                  primer: TCACACAGGAAACAGCTATGAC
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/organism≃"Medicago truncatula"
/db_xref="taxon:3880"
                                                           ocation/Qualifiers
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Pred. No. 2.1e+03;
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1 tgttggannnnnnnntttgata
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similar to
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: L. Jeffrey Mei
Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tumor Gene Index
Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostom1; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 298)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AA515145.1
                                                                                                                                                                                                                                                                                                                                                                                                                      CDNA Library Preparation: David B. Krizman, Ph.D.

CDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing Center

Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:

www-bio.llnl.gov/bbrp/image/image.html

seq.primer: -40ml3 fwd. ET from Amersham
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      National Cancer Institute, Cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            nullian
                                                                      Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                          quality sequence stop: 295.
                                                                                                                                                             76
                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /clone="NF082F04DT"
/clone_lib="Drought"
/clone_lib="Drought"
/tissue_type="Plantlets"
/tissue_type="Pooled timepoints"
/dev_stage="Pooled timepoints"
/note="Vector: Lambda Zap; Contains a mixture of entire
plantlets harvested in a series of days-post-watering
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      timepoints.
76 c
                                                                                                                                                                                           /note="Vector: pAMP10; mRNA made from liposarcoma, cDNA made by oliqo-dT priming. Non- directionally cloned. Size-selected on agarose gel, average insert size 600 bp. Reference: Krizman et al. (1996) Cancer Research
                                                                                                                                                             56:5380-5383.
63 c
                                                                                                                                                                                                                                                                                                                     /db_xref="taxon:9606"
/clone="IMAGE:939936"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               298 bp mRNA EST 20-AUG-1997
NCI_CGAP_Lip2 Homo sapiens cDNA clone IMAGE:939936
contains Alu repetitiive element;contains element MSR1
element; mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                      /organism="Homo sapiens"
                                                                                                                                                                                                                                                                 /lab_host="DH10B"
                                                                                                                                                                                                                                                                                  /tissue_type="liposarcoma"
                                                                                                                                                                                                                                                                                                      /clone_lib="NCI_CGAP_Lip2"
                                                                                                                                                                                                                                                                                                                                                                                      ocation/Qualifiers
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Pred. No. 2.2e+03;
0; Mismatches 8
                      22
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                                                                        Score 14; D
Pred. No. 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Jeffrey Medeiros, M.D., Michael
                                                                                                                                                             91 g
                                                         Mismatches
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                                                                                                                                                                 83
                                                                                          DB 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Genome
                                                                          2e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Anatomy Project (CGAP),
                                                                                          Length 298;
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                                                           Indels
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TGTTGGAGAATGACATTTGATA

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RESULT 13
AZ037213
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                                        DEFINITION
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ACCESSION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         source
                                                                                                                                                        157 TGTTGGATTGATGGGTTTGATA
                                                                                                                                                                                                                                                        Local Similarity
                                                                                                                                                                               1 tgttggannnnnnnntttgata
                                    AZ037213 336 bp DNA
RPCI-23-364L15.TV RPCI-23 M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     http://chrysie.tamu.edu/medicago
Seg primer: SKmod (CTA gAA CTA gi
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Texas A&M University: T262522e
TIGR sequence name: MTIAE96TK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Texas A&M University
CTOLlege Station, TX 77843-3258,
Tel: 409 845 7707
Fax: 409 845 2891
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosidae; eurosids I; Fabales; Fabaceae; Papillonoideae; Trifolieae;
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EST402778 KV1
  AZ037213
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Department of Biology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: VandenBosch K
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Fraser, C.M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Town, C.D., Bowman, C.L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             VandenBosch,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Medicago truncatula
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                    DNA sequence.
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                                                                                                                                                                                                                                                                                                                                                     95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                from roots of Medicago truncatula 24 hours after inoculation Sinorhizobium meliloti \,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           kate@mail.bio.tamu.edu
                                                                                                                                                                                                                                                                                                                                             /note="Vector: pBluescript SK -; Site_1: EcoRI; Site_2: XhoI; CDNA was prepared from polyA+ enriched RNA. The cDNA was directionally ligated into the Unizap XR vector from Stratagene and packaged using Gigapack III Gold packaging extracts. Plasmids containing cDNA inserts were excised from the recombinant lambda-Zap phage using Ex-assist helper phage and propagated in XLOLR cells."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /organism="Medicago truncatula"

/cultivar="genotype A17"

/db_xref="taxon:3880"

/clone="pkVl-3023"

/clone_lib="KVl"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /lab_host="E. coli strain XLOLR"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /tissue_type="Seedling roots"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /dev_stage="24 hours post-inoculation with Sinorhizobium
                                                                                                                                                                                                                                                      63.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Medicago truncatula cDNA clone pKV1-3023, mRNA
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an,C.L., Craven,M.B., Hansen,T.S., Holt,I.E.
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                                                                                                                                                        178
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                                    Mus
                                  musculus genomic clone RPCI-23-364L15
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2.2e+03;
ches 8;
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RESULT 14
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LOCUS
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AUTHORS
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                   AUTHORS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Clones are derived from the mouse BAC library RPCI-23. For BAC library availability, please contact Pieter de Jong (pieter@dejong.med.buffalo.edu). Clones may be purchased from BACPAC Resources (http://bacpac.med.buffalo.edu/orderingframe.htm) or from Resea ch Genetics (info@resgen.com). BAC end page: http://www.tigr.org/tdb/bac_ends/mouse/bac_end_intro.html plate: 364 row: L column: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mus musculus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleosto
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
                                   Eukaryota; Metazoa;
Mammalia; Eutheria;
1 (bases 1 to 342)
                                                                                                                                                                                                        BH083614 342 bp DNA
RPCI-24-293E7.TJ RPCI-24 Mus
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Other_GSSs: RPCI-23-364L15.TJ
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GSS.
                                                                                                                                                                     DNA sequence
BH083614
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Contact: Shaying Zhao
Department of Eukaryot
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   and Fraser, C.M. Mouse BAC End S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     zhao,S., Nierman,W., Feldblyum,T., Malek,J., Shatsman,S.,
,B., Levins,M., Mcgann,S., Tsegaye,G., Geer,K., Krol,M.,
               Zhao, S., Nierman, W.,
                                                                                          Mus musculus
                                                                                                                                                     BH083614.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Seq primer: T7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tel: 301 838 0200 Fax: 301 838 0208
Tsegaye,G.,
                                                                                                                house mouse.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Email: szhao@tigr.org
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                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 97
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note-"Organ: Kidney/Brain; Vector: pBACe3.6; Site_1; EcoRI; Site_2: EcoRI; Female C57BL/6J mouse kidney and/or brain genomic DNA was isolated and partially digested with a combination of EcoRI and EcoRI Methylase. Size selected DNA was cloned into the pBACe3.6 vector at the EcoRI sites. The ligation products were transformed into DH10B electrocompetent cells (BRL Life Technologies). " a 37 c 45 g 157 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /db_xref="taxon:10090"
/clone="RPCI-23-364L15"
/clone_lib="RPCI-23"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /organism="Mus musculus"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /sex="Female"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ocation/Qualifiers
 Geer, K.,
                                                                                                                                                     GI:14903211
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                                                                        Chordata;
                                                         Rodentia;
                                                                                                                                                                                                                                                                                                                                                            22
Krol, M., Shvartsbeyn, A.,
                 Malek,J.,
                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                                                     Score 14;
Pred. No.
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                                                                                                                                                                                                        musculus genomic clone RPCI-24-293E7,
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                                                     Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
                   Shatsman, S.,
                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 13;
                                                                                                                                                                                                                                                                                                                                                                                                 .2e+03;
les 8;
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                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 336;
                   Akinret, B.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Shatsman, S.,
 Gebregeorgis, E.,
                                                                                                                                                                                                                                18-JUL-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Department of Eukaryotic Genomics The Institute for Genomic Research 9712 Medical Center Dr., Rockville, Tel: 301 838 0200 Fax: 301 838 0208
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Russell,D., de Jong,P. and Fraser,C.M.
Mouse BAC End Sequences from Library RPCI-24
Unpublished (1999)
Contact: Shaying Zhao
                                                                                                                                                                                                                                              Caenorhabditis elegans.
Caenorhabditis elegans
Eukaryota; Metazoa; Nematoda; Chromadorea; Rhabditida; Rhabditoidea; Rhabditidae; Peloderinae; Caenorhabditis.
1 (bases 1 to 349)
1 (bases 1 to 349)
1 (bases 1, Shin-1, T., Thierry-Mieg, J., Thierry-Mieg, D., Suzuki, Y. and Sugano, S.
                                                                                                                                                                                                                                                                                                                                                                                                                         AU112918 349 bp mRNA EST 19-OCT-2000 AU112918 unpublished oligb-capped cDNA library Caenorhabditis elegans cDNA clone yk/148e6 5', mRNA sequence.
                                                                                                                                                                                      A complementary view of the C. Unpublished (2000)
Contact: Yuji Kohara
                                                                      Genome Biology Lab.
National Institute of Genetics.
Yata 1111, Mishima, Shizuoka 411,
Tel: 81-559-81-6854
Fax: 81-559-81-6855
Email: ykohara@lab.nig.ac.jp.
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                                                    ykohara@lab.nig.ac.jp.
Location/Qualifiers
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/note="Vector: pTARBAC1; Site_1: BamH1; Sites using MboI partially digested male C57BL/6J
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155 c 81 g 115 t
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Search completed: December 15, 2001, 02:33:48 Job time: 4988 sec

Page 9

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	Description U60594 Leptospira AB007014 Spirochae AF003953 Leptospira U94976 Leptospira U94976 Leptospira U94976 Leptospira U94977 Leptospira U94977 Leptospira U94979 Leptospira U046176 Homo sapi AC0261779 Nephrosel AF137379 Nephrosel AF137379 Nephrosel AF137379 Nephrosel AF220067 Drosophil AC015209 Drosophil AC015209 Drosophil AC015209 Drosophil AC026327 Homo sapi AC026327 Homo sapi AC026327 Homo sapi AC026327 Homo sapi AC026315 Homo sapi AC02315 Homo sapi AC035742 Homo sapi AC02315 Homo sapi AC035742 Homo sapi AC04363 Canorhabdi AL04365 Arabidops	

RESULT 1
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LEPGOSPIRA fainei 16S ribosomal RNA gene, partial sequence.
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UGOS94.1 GI:1408219
KEYWORDS
SOURCE
LEPTOSPIRA fainei
Bacteria; Spirochaetales; Leptospiraceae; Leptospira.
REFERENCE
AUTHORS
BILLINGPURST, M.L., Letocart, M., Merien, F. and Serrano, M.S.
FITLE
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MEDLINE
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REFERENCE
AUTHORS
BILLINGPURST, Bacteriol. 48 Pt 3, 851-858 (1998)
RIBERENCE
AUTHORS
BILLINGPURST, M.L., Letocart, M., Merien, F., Serrano, M.S.
BILLINGPURST, Bacteriol. 48 Pt 3, 851-858 (1998)
BILLINGPURST, M.C., Letocart, M., Merien, F., Serrano, M.S.
BILLINGPURST, Bacteriol. 48 Pt 3, 851-858 (1998)
Perolat, P.
BILLINGPURST, M.L., Letocart, M., Merien, F., Serrano, M.S. and
Perolat, P.

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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                                                                                                                                                                                                                                                                                                                                                                                        Submitted (05-SEP-1997) to the DDBJ/EMBL/GenBank databases. Mi-Su Shin, Laboratory of Marine Wolecular Microbiology, Faculty of Agriculture, Kyoto University; Olwakecho, Kitashirakawa, Sakyoku, Kyoto, Kyoto 306-01, Japan (E-mail:misun@kais.kyoto-u.ac.jp, Tel:075-753-6224, Fax:075-753-6226)
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Bacteria; Spirochaetales;
1 (bases 1 to 354)
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strain:FO-95) DNA.
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Shin, M., Y
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Best Local
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                                                                                       source
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AF003953
AF003953.1
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U94974
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Woo,T.H.S., Patel,B.K.C.,
Norris,M. and Dohnt,M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bacteria; Spirochaetales; Leptospira
1 (bases 1 to 283)
WOO,T.H.S., Patel,B.K.C., Cinco,M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20;
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                                                                                                                                                                                                Bacteria; Spirochaetales; Leptospiraceae; Leptospira.

1 (bases 1 to 288)
Woo,T.H.S., Smythe,L.D., Symonds,M., Norris,M., Dohnt,M.,
Brenner,D.J. and Patel,B.K.C.
Identification of Leptospira inadai by continuously monit
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Woo,T.H.S., Smythe,L.D., Symonds,M.,
Brenner,D.J. and Patel,B.K.C.
Direct Submission
                                                                                                                                                                                                                                                                                                                  U94974.1
                                                                                                                 Submitted (24-MAR-1997) School Brisbane, QLD 4111, Australia
                                                                                                                                                                                      fluorescence during 
Unpublished
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Leptospira inadai
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60 c 88 9
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/strain="48/95"
                                       /db_xref="taxon:29506"
-<1. .>288
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             /product="16S
61 c
                                                               /organism="Leptospira inadai"
/strain="79/95"
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Pred. No. 78;
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               ribosomal RNA<sup>®</sup>
88 g ·62 t
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Query Match

85.5%;

Score

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JOURNAL
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Best Local Similarity 90.1
Matches 20; Conservative
                                                                                                                    TITLE
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Matches 20; Conservative
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U94976
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U94975
                                                                                                    Bacteria; Spirochaetales; Leptospiraceae; Leptospira.

1 (bases 1 to 288)

Moo,T.H.S., Smythe,L.D., Symonds,M., Norris,M., Dohnt,M.,
Brenner,D.J. and Patel,B.K.C.

Identification of Leptospira inadai by continuously monit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WOO,T.H.S., Smythe,L.D., Symonds,M., Norris,M., Dohnt,M., Brenner,D.J. and Patel,B.K.C. Identification of Leptospira inadai by continuously monit fluorescence during rapid cycle PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2 (bases 1 to 288)
Woo,T.H.S., Smythe,L.D., Symonds,M.,
Brenner,D.J. and Patel,B.K.C.
Submitted (24-MAR-1997) School of
Brisbane, QLD 4111, Australia
Location/Qualifiers
                                       2 (bases 1 to 288)
WOO,T.H.S., Smythe,L.D., Symonds,M.,
Brenner,D.J. and Patel,B.K.C.
Direct Submission
                                                                                                                                                                                                                          U94976.1
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Leptospira inadai
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                                                                                        Unpublished
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/strain="68/94"
/db_xref="taxon:29506"
<1. >288
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61 c 88 g 62 t
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90.9%;
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Pred. No. 78;
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0; .Mismatches
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                                                                 Norris, M.,
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                           Griffith University,
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                                                               Dohnt, M.
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VERSION
KEYWORDS
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ORIGIN
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AUTHORS
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Matches 20
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Best Local Similarity
Matches 20; Conserv
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U94978
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U94977
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1 (bases 1 to 288)

Woo,T.H.S., Smythe,L.D., Symonds,M., Norris,M., Dohnt,M.,
Brenner,D.J. and Patel,B.K.C.

Identification of Leptospira inadai by continuously monit.

fluorescence during rapid cycle PCR
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20; Conserv
         Leptospira inadai.
Leptospira inadai
Bacteria; Spirochaetales; Leptospiraceae; Leptospira.
1 (bases 1 to 288)
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Woo,T.H.S., Smythe,L.D., Symonds,M.,
Brenner,D.J. and Patel,B.K.C.
Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                     Submitted (24-MAR-1997) School Brisbane, QLD 4111, Australia
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 WOO, T.H.S.,
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/strain="346/95"
/db_xref="taxon:29506"
<1. .>288
                                                                                                                                                                                                                                                                                                                                         /organism="Leptospira inadai"
/strain="268/95"
/db_xref="taxon:29506"
<1..>288
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61 c 88 g 62 t
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61 c 88 g 62 t
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                                                                             GI:2735450
 Smythe,L.D.,
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90.9%;
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Pred. No. 78;
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Pred. No. 78;
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Symonds, M., Norris, M., Dohnt, M.,
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Griffith University,

Dohnt, M.,

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partial sequence.

01-JAN-1998

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Leptospira inadai 16S ribosomal RNA gene,
U94979
U94979.1 GI:2735/53
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Woo,T.H.S., Smythe,L.D., Symonds,M., Norris
Brenner,D.J. and Patel,B.K.C.
Direct Submission
Submitted (24-MAR-1997) School of Science,
Brisbane, QLD 4111, Australia
Location/Qualifiers
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Identification of Leptospira inadai by
                                                20;
                                                                                                                                                                                                       Submitted (24-MAR-1997) School
Brisbane, QLD 4111, Australia
Location/Qualifiers
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Identification of Leptospira inadai by continuously monitoring fluorescence during rapid cycle PCR
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/strain="218/95"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Submitted (05-SEP-1997) to the DDBJ/EMBL/GenBank databases. Mi-Sun Shin, Laboratory of Marine Molecular Microbiology, Faculty of Agriculture, Kyoto University; Oiwakecho, Kitashirakawa, Sakyoku, Kyoto, Kyoto 306-01, Japan (E-mail:misun@kais.kyoto-u.ac.jp, Tel:075-753-6224, Fax:075-753-6226)
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Spirochaeta sp. 16S r
AB007012
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Spirochaeta sp.
Bacteria; Spirochaetales; Spirochaetaceae;
[] (bases 1 to 353)
                                                                                                                                                                                                                       Arabidopsis thaliana cDNA to mRNA.

Arabidopsis thaliana

Arabidopsis thaliana

Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;

Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;

Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.

1 (bases I to 1012)
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16S ribosomal RNA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Direct Submission
                                                                                                                                                                          RNS2: a senescence-associated RNase of Arabidopsis that diverged
                                                                                                                                                                                                        Taylor,C.B., Bariola,P.A., delCardayre,S.B.,
                                                                                                                                             from the S-RNases before speciation Proc. Natl. Acad. Sci. U.S.A. 90 (11),
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80 c 108 g
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                                                           /organism="Arabidopsis
/db_xref="taxon:3702"
                                                                                                           Location/Qualifiers
'gene="RNS2"
                                 'gene="RNS2"
                                                                                                                                                                                                                                                                                                                                      GI:289209
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Arabidopsis genome 
Unpublished
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1 (bases 1 to 9423)
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AJ275979.1 GI:13509286
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/protein_id="AA55406.1"
/protein_id="AA55406.1"
/db_xref="Gi:289210"
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/CRGTRHCCSKNACCRGSDAPTQFTIHCLWPDYNDGSWPSCCYRSDFKEKEISTLMDG
LEKYWPSLSCGSPSSCNGGKGSFWGHEWEKHGTCSSPVFHDEYNYFLTTLNLYLKHNV
TDVLYQAGYVASNSEKYPLGGIYTAIQNAFHITPEVVCKRDAIDEIRICFYKDFKPRD
CVGSQDLTSRKSCPKYVSLPEYTPLDGEAMVLKMPTEREAL"
                                                                                                                                  /translation="MKALILYGGFGTTLRPLTLSFPKPLVDFANKPMILHQIEALKAV GVDEVVLAINYQDFEVMLNFLKDDETKLEJKITGCGGTEDLGTAGLALARDKILDGSG EPFFVLMSDV15EYPLKELKDFEKKHGGEASLAVTKVDEPSKYGVVYMSESTGRVEKF VEKPKLYVGNKINAGIYLLNPSVLDKIELRPTSIEKETFPKIAAAQGHYAMVLPGFWM DIGQPRDYITGLGLYLDSFKKSSPAKLTSGPHIVGNVLVDETAATIGEGCLIGPDVAIG PGCIVESGVRLASGVRLXKHAGISSSIIGWHSTVGQWARIENWTILGEDVHVS DEIYSNGGVVLPHKEIKSNILKPEIYM"
          1597. .1659
/number-21
                                     /number=1
1597. .169
                                                                                                                                                                                                                                                                        /product="GDP-mannose pyrophosphorylase"
/protein_id="CAC35355.1"
/db_xref="GI:13509287"
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90.9%; Pred. No. 76;
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1 (bases 1 to 72590)

1 (Lin,X., Kaull,S., Rounsley,S.D., Shea,T.P., Benito,M.-I., Town,C.D., Fujii,C.Y., Mason,T.M., Bowman,C.L., Barnstead,M.E., Fulli,C.Y., Mason,T.M., Bowman,C.L., Barnstead,M.E., Ronning,C.M., Feldblyum,T.V., Buell,C.R., Ketchum,K.A., Lee,J.J., Ronning,C.M., Feldblyum,T.V., Buell,C.R., Ketchum,K.A., Lee,J.J., Ronning,C.M., Koo,H., Moffat,K.S., Cronin,L.A., Shen,M., Vanaken,S.E., Umayam,L., Tallon,L.J., Gill,J.E., Adams,M.D., Carrera,A.J., Creasy,T.H., Goodman,H.M., Somerville,C.R., Copenhaver,G.P., Preuss,D., Wlerman,W.C., White,O., Eisen,J.A., Salzberg,S.L., Fraser,C.M. and Venter, J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20;
             Direct Submission
Submitted (09-MAR-2000) The Institute for Genomic Research, 9712
Medical Center Dr., Rockville, MD 20850, USA
On Dec 17, 1999 this sequence version replaced 91:2642152.
The sequence and annotation of chromosome 2 were merged from those of the individual clones on this chromosome after removing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AC003000 72590 bp DNA PLN Arabidopsis thaliana chromosome II section 214 of complete sequence. Sequence from clones T517. AC003000 AE002093 AC003000.2 GI:6598383
overlaps. For detailed information, please see the TIGR web site
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                                                                                                                                                                                                                Nature 402 (6763),
20083487
                                                                                                                                                                                                                                                                                                      Venter,J.C.
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2099. .27
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/protein_id="CAC35357.1"
/db_xref="GI:13509289"
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1693 c 1780 g 3155 t
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/product="vacuolar ribonuclease"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     F6P23, F5J6, T17A5, and T13L16, the ESSA group for sequencing clone F13D4, and Scott Jackson, Jining Jiang, Klaus Meyer, Eric Richards and Satoshi Tabata for helpful assistance. In addition, we would like to thank the TIGR Bioinformatics Department, especially Lixin Zhou, Hanif Khalak, Michael E. Heaney, Lily Fu, Feng Liang, Jeremy Peterson, Michael Holmes, and Delwood Richardson for software and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  http://genome.wustl.edu/eddy/tRNAscan-SE/). Simple repeats were identified by repeatmasker (Arian Smit, http://ftp.genome.washington.edu/RM/RepeatMasker.html). Genes are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (ftp://arthur.epm.ornl.gov/pub/xgrail), Genefinder (Phil Green, University of Washington), Genscan (Chris Burge, http://gnomic.stanford.edu/GENSCANW.html), and NetPlantGene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Genes were identified by a combination of three methods: Gene prediction programs including GRAIL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        predicted by tRNAscan-SE (Sean Eddy,
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complement(2759...2862)
/rpt_family="POLY_A"
complement(4584...>5909)
/gene="Att939710"
complement/
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /HOTE="T517.1; Similar to GP|2245012|gnl|PID|e327000|297341" complement(4584, .5909) gene="At239710" /note=""."
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complement(1266. .1291)
/rpt_fam.')
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/gene="At2g39710"
/note="T517.1; similar to
                                                                                                                                                                                                                                                                                                      /codon_start=1
/protein_id="AABB7120.1"
/protein_id="AABB7120.1"
/db_xref="GI:2642153"
/translation="MDSSSSSSLSLSKNFLRISVLLLIFPLTFCKTSSTNQTLLFSLK
/translation="MDSSSSSSLSLSKNFLRISVLLLIFPLTFCKTSSTNQTLLFSLK
TQKLPQSSSDKLSFRHNVTLTVTLAVGDPPQNISMVLDTGSELSWLHCKKSPNLGSVF
TQKLPQSSSTYSPVPCSSPICRTRTRDLPIPASCDPKTHLCHVAISYADATSIEGNLAHET
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complement(1000. .102
                                    SDLLGIEAFVIGHHHQQNVWMEFDLAKSRVGFAGNVRCDLASQRLGLRD'complement(<10039. .11304)
                                                                                                                       SVFVPDHTGAGQTMVDSGTQFTFLMGPVYTALKNEF1TQTKSVLRLVDDPDFVFQGTM
DLCYKVGSTTRPNFSGLPMVSLMFRGAEMSVSGQKLLYRVNGAGSEGKEEVYCFTFGN
                                                                                                                                                                                                              FVIGSVTRPGTLFGCMDSGLSSNSEEDAKSTGLMGMNRGSLSFVNQLGFSKFSYCISG
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/chromosome="II"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note="unknown protein"
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//POCULITIES AAB87121.1"
//db_xref="G1:2642154"
//db_xref="G1:26421
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      complement(<10039. .11304)
/gene="At2g39720"
/note="T517.2"</pre>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /db_xref="GI:2642170"
/translation="MAAAVSTGAINRAPLSLNGSGSGAVSAPASTFLGKKVVTVSRFAOSHKKSNGSEKVLAVEDKOTDGDRWRGLAYDTSDDQDDITRCKGMVDSVFQAPMGTGONKKSNGSEKVLAVEDKOTDGDRWRGLAYDTSDDQDDITRCKGMVDSVFQAPMGTGTHAVLSSYEVSQGLRQYNLDNMMDGFYIAFAFMDKLVVHITKNFLTLPNIKVPLIGIWGGKGQCKSFQCELVMAKGINFLMMSAGELESGAMGEPAKLIRORYREAADLIK LGIWGGKGQCKSFQCELVMAKGINPLMMSAGELESGAMGEPAKLIRORYREAADLIK LGIWGGKGQCKSFQCELVMAKGINPLMMSAGELESGAMGEPAKLIRORYREAADLIK LGIWGCKGIRTDKIKDEDIVTLV LGIWGCKGCIFTDKIKDEDIVTLV LGIWGCKGGIRTDKIKDEDIVTLV DQFPGQSIDFFGALRARVYDDEVRKFVESLGVEKIGKBLUNGSEPPVFEQPEMTYEK DQFPGQSIDFFGALRARVYDDEVRKFVESLGVEKIGKBLUNGSEPPVFEQPEMTYEK
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complement(join(13484...13599,13724...14005,14087...14559.

14651...14765,14836...14941,15033...15349,15834...15878))

/gene="At2g39730"

/note="atternative splice form 1 GP}166834|M86720"

/codon_start=1
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/gene="At2g39730"
/note="T517.3"
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/rpt_family="POLY_A"
complement()oin(13323. .13599,13724. .14005,14087. .14559,
14651. .14736,14836. .14941,15033. .15349,15834. .16055))
/gene="At2g39730"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              complement(join(13579. .13599,13713. .14005,14087. .14559)14651. .14736,14836. .14941,15033. .15349,15834. .15878))/gene="at2339730"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DQFPGQSIDFFGALRARVYDDEVRKFVESLGVEKIGKRLVNSREGPPVFEQPEMTYEK
LMEYGNMLVMEQENYKRVQLAETYLSQAALGDANADAIGRGTFYGKGAQQVNLPVPEG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        KGKMCCLFINDLDAGAGRMGGTTQYTVNNQMVNATLMNIADNPTNVQLPGMYNKEENA
RVPIICTGNDFSTLYAPLIRDGRMEKFYWAPTREDRIGVCKGIFRTDKIKDEDIVTLV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /translation="MAAAVSTYGAINKAPLSLNGSGSGAVSAPASTFLGKKVYTYSRF
AQSNKKSNGSFKVLKEDKQTDGDRWRGLAYDTSDDQDITRGKGWVDSVFQAPMGT
GTHHAVLSSYEYVSQGLKQYNLDNWMDGFYIAPAFMDKLVVHITKNFLTLPNIKVPLI
LGIWGGKGQGKSFQCELVMAKMGINPIMMSAGELESGNAGEPAKLIRQRYREAADLIK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /product="Rubisco activase"
/protein_id="AAB87122.1"
/db_xref="GI:2642155"
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/codon_start=1
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                                                                                                                                                                                                                                                            join(<18783. .18921,19188. .19352,19496. .
19965. .20034,20131. .20261,20593. .>21237
                                                                                                                                                                                                                                                                                                                                                                                                                                      LMEYGNMLVMEQENVKRVQLAETYLSQAALGDANADAIGRGTFYGKTEEKEPSK"
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/gene="At2g39740"
/gene="7517.4; predicted by genscan"
/note="7517.4; predicted by genscan"
join(18783. .18921,19188. .19352,19496. .19674,19766. .19861,
19965. .20034,20131. .20261,20593. .21237)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /product="Rubsico activase"
/protein_id="AAB87137.1"
                                                                                                                                                                                                                        /gene="At2g39740"
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JOURNAL REFERENCE
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VERSION
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AF165146
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Biotechnology, Beutenbergstrasse 11, Jena 07745, Germany On Jun 7, 2001 this sequence version replaced gi:8151945
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2 (bases 1 to 121524).
Schudy, A., Blechschmidt, K., Schillhabel, M., Baumgart, C., Menzel, U., Weber, J., Schattevoy, R. and Rosenthal, A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Lehmann, R., Menzel, U., Polley, A., Siddiqui, R., Taudien, S., Wen, G., S Rosenthal, A. and Platzer, M. Chromosome 8 genomic sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AF165146 121524 bp DNA HTG 07-JUN HOMO Sapiens chromosome 8 clone CTA-397H3 map 8q12-8q13, SEQUENCING IN PROGRESS ***, 7 unordered pieces.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 (bases 1 to 121524)
Schilhabel, M.B., Baumg
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HTG; HTGS_PHASE1; HTGS_FULLTOP.
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NOTE: This is a 'working draft' sequence. It currently
                                                                         Sequencing vector. M13; 100% of reads Chemistry: Dye-terminator Big Dye; 100% of reads Assembly program: Phrap; version 0.990329 Consensus quality: 115981 bases at least 040 Consensus quality: 119312 bases at least 020 Quality coverage: 9.57 in Q20 bases; sum-of-continuation of the control 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Center: Insitute of Molecular Biotechnoloy Center code: IMB

Web site: http://genome.imb-jena.de/
Contact: gscj-submit@genome.imb-jena.de

Project Information
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Center clone name: CTA-397H3
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22328. 22585,22834. 22993,23212. 23322,23481. 23751,
23847. 23996,24089. >>24814))
complement(21380. >24814)
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LRGATVQPFCSFVSNLFTRWGDLDISVDLFSGSSILFTGKKQKQTLLGHLLRALRASG
LWYKLQFVLHARVPILKVVSGHQRISCDISIDLDGLLKSRFLFWISSIDGRRBDLVL
LVKEWAKAHNINDSKTGTFNSYSLSLLVIFHFQTCVPAILPPLRVIYPKSAVDDLTGV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /gene="At2g39740"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (06-JUL-1999) Genome Analysis, Institute of Molecular Logy, Beutenbergstrasse 11, Jena 07745, Germany
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SBirren, B., Linton, L., Nusbaum, C., Lander, E., Abraham, H., Allen, N., Anderson, S., Barna, N., Bastien, V., Beda, F., Boguslavkiy, L., Anderson, S., Barna, N., Burkett, G., Campopiano, A., Castle, A., Choepel, Y., Colangelo, M., Collins, S., Collymore, A., Cooke, P., DeArellano, K., Dewar, K., Diaz, J.S., Dodge, S., Ferreira, P., FitzHugh, W., Gage, D., Galagan, J., Gardyna, S., Ginde, S., Goyette, M., Graham, L., Grand-Pierre, N., Hagos, B., Heaford, A., Horton, L., Iliev, I., Johnson, R., Jones, C., Kann, L., Karatas, A., LaRocque, K., Lamazares, R., Landers, T., Lehoczky, J., Levine, R., Lieu, C., Liu, G., Macdonald, P., Marquis, N., McCarthy, M., McEwan, P., McKernan, K., Morrow, J., Murphy, T., Naylor, J., Norman, C.H., O'Conmor, T., O'Donnell, P., O'Neil, D., Olivar, T.M., Oliver, J., Peterson, K., Pierre, N., Pisani, C., Pollara, V., Raymond, C., Rieback, M., Riley, R., Rogov, P., Rothman, D., Roy, A., Santos, R., Schauer, S., Severy, P., Sougnez, C., Spencer, B., Stange-Thomann, N., Stojanovic, N.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Homo sapiens chromosome 8 clone RP11-22E14 map 8,
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17: gap of unknown length
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Submitted (08-OCT-2000) whitehead Institute/MIT Center for Research, 320 Charles Street, Cambridge, MA 02141, USA On Feb 15, 2001 this sequence version replaced gi:10717228
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              http://ftp.genome.washington.edu/RM/RepeatMasker.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             All repeats were identified using Rosmit, A.F.A. & Green, P. (1996-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NOTE: This is a 'working draft' sequence. It currently consists of 67 contigs. The true order of the pieces is not known and their order in this sequence record is arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown. This record will be updated with the finished sequence.
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Center clone name: 22_E_14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  web site: http://www-seq.wi.mit.edu
Contact: sequence_submissions@genome.wi.mit.edu
------ Project Information
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                                                                                                                                                                                                                                                                                                                     56 59965: gap of 100 bp
66 62260: contig of 2295 bp
66 62260: contig of 100 bp
61 62360: gap of 1760 bp
61 64120: gap of 100 bp
61 64220: gap of 100 bp
62 6979: contig of 1759 bp
61 6679: gap of 67 6733 bp
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51438: gap of 2005

53743: contig of 2305

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38963: contig of 1827 bp
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33540: contig of 858
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92728: contig of 4907 bp in
2828: gap of 100 bp
96552: contig of 3824 bp in
752: gap of 100 hr
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55260: contig of 1417 bp
5360: gap of 100 bp
57843: contig of 2483 bp
7943: gap of 100 bp
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31038: contig of 1427 l
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29511: contig of 830 bp
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26183: contig of 1192 bp in length
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51338: contig of
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96: gap of 100 bg
44598: contig of 1602
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103011 107188: contig of 4178 bp in length
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111048 111147: gap of 100 bp
111048 1116741: contig of 5594 bp in length
116742 116841: gap of 100 bp
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Copyright (c) 1993 - 2000 Compugen Ltd
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S. epidermidis ope Human olfactory re Adenovirus pACTSG2 Adenovirus SCAR.RG C. elegans inhibit	H. pylori GHPO 131 H. man hypoxia resp EST clone HK650. Human olfactory re	olfac secre secre dopsis	Maize acetyl CoA c Maize ACCase enzym Maize acetyl CoA c Potato pollen cell	etyl-CoA-ca ize acetyl oRI fragmen leprae gyr maize ACCa	Maize male fertill Eucalyptus grandis Human colon cancer Human secreted pro Human secreted pro Human secreted pro Human colon cancer Arabidopsis thalia Arabidopsis thalia

ALIGNMENTS

AAV58901

AAV58901 standard;

DNA; 22 ВР

Leptospira rRNA gene nucleotide sequence

20-JAN-1999 AAV58901;

(first entry)

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New isolated pathogenic Leptospira bacterium - useful for, edeveloping products for conferring protective immunity, and prophylactic or therapeutic treatment
                                                                                                    07-MAR-1997;
                                                                                                                                                                                         Infection; pathogenic Leptospira; protective immunity; therapy;
diagnosis; ss.
                                        WPI; 1998-520791/44.
                                                        Chappel RJ;
                                                                         (AGRI-) AGRIC VICTORIA SERVICES PTY LTD. (PIGR-) PIG RES & DEV CORP.
                                                                                                                      06-MAR-1998;
                                                                                                                                      17-SEP-1998.
                                                                                                                                                        WO9840099-A1
                                                                                                                                                                        Leptospira sp.
                                                                                                    97AU-0005494.
                                                                                                                      98WO-AU00145
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RESULT
AAV58897
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Query Match
Best Local Similarity
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This sequence represents a Leptospira DNA sequence isolated from the pathogenic Leptospira (LS) bacterium of the invention. The bacterium belongs to serogroup Hurstbridge of servovar hurstbridge or the species L. fainel. The LS bacteria can be used for conferring protective immunity against pathogenic LS bacteria in humans or animals. The bacteria can also be used for prophylactic or therapeutic treatment of infections. The DNAs and antibodies may also be used for detection and diagnosis of past or present LS infection.
                                                                              pathogenic Leptospira (LS) bacterium of the invention. The bacterium belongs to serogroup Hurstbridge or serovar hurstbridge or the species L. fainei. The LS bacteria can be used for conferring protective immunity against pathogenic LS bacteria in humans or animals. The bacteria can also be used for prophylactic or therapeutic treatment of infections. The DNAs and antibodies may also be used for detection and
                                                                                                                                                                                               New isolated pathogenic Leptospira bacterium - useful for, developing products for conferring protective immunity, an prophylactic or therapeutic treatment
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                                               Sequence
                                                                                                                                                   This sequence represents a Leptospira DNA sequence isolated from the
                                                                                                                                                                                                                                                                                       (AGRI-)
(PIGR-)
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                                                                  infections. The DNAs and antiboodiagnosis of past or present LS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAV58897 standard; DNA;
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                                               22
                                                                                                                                                                         Page 70; 94pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Page
                                                                                                                                                                                                                                                                                                                                                                                                                                          pathogenic Leptospira; protective immunity; therapy;
ss.
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                                               В₽;
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 No.;
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           Length 22;
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RESULT AAA99365

4

AAA99365 standard; DNA; 1012

ВP

DE XX AC XX

22-JAN-2001 Plant PrAG1

promoter related gene sequence.

(first entry)

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RESULT
AAV58896
ID AAV5
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                                                                                                                                  Query Match
Best Local
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(PIGR-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New isolated pathogenic Leptospira bacterium - useful for, edeveloping products for conferring protective immunity, and prophylactic or therapeutic treatment
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                                                                                                                                                                                                                                       Sequence 1477 BP; 390 A; 334 C;
                                                                                                                                                                                                                                                                                        diagnosis of past or present LS infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 15; Page 69-70;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1998-520791/44.
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tgttggatcacaagatttgata 175
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                                                                                                          Similarity
22; Conserv
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                                                                                                            Conservative
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                                                                                                                                     100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        94pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Leptospira; protective immunity; therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1477
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                                                                                                                                     Score 22;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           YTG
                                                                                                                                                                                                                                       439 G;
                                                                                                            Mismatches
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                                                                                                                                                              DВ
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RESULT
AAC37325
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This invention relates to a novel plant promoter gene. The promoter is located in plant reproductive tissue, and the invention includes transgenic plants containing the promoter. The promoter can be used to produce plants which have a diminished reproductive capacity or which a sterile. The constructs can also be used to transform agronomically important plants in which modulation of reproductive capacity (particularly the timing and abundance of flowering) is desirable, e.g. cereals, rice, maize, wheat, barley, oats, rye, soybean and canola The present sequence represents DNA encoding the plant reproductive promoter (PrAGI) of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel
which
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                                                                                                                                             Hybridisation assay; genetic mapping; gene expression control;
protein identification; signal transduction pathway;
                                                                                                                                                                                                                                                             AAC37325 standard; DNA; 1069
                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 1012 BP; 287 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim
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P-PSDB; AAB26796.
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(TASM-) TASMAN BIOTECHNOLOGY LTD.
(UNMT ) UNIV MICHIGAN TECHNOLOGICAL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17-MAR-1999;
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                           25-FEB-2000; 2000EP-0301439
                                                      06-SEP-2000.
                                                                                                     Arabidopsis thaliana.
                                                                                                                              metabolic pathway; promoter; termination sequence;
                                                                                                                                                                                   Arabidopsis thaliana DNA fragment SEQ
                                                                                                                                                                                                            17-OCT-2000
                                                                                                                                                                                                                                     AAC37325;
  25-FEB-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            plant reproductive tissue promoter, useful to produce plants have a diminished reproductive capacity or which are sterile
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GK, Liu J,
                                                                                                                                                                                                                                                                                                                                                                                 20;
                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Page 42-43;
                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                 Conservative
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  99US-0121825.
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Pred. No. 5.8;
O; Mismatches
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8
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                                                                                                                                                                                                                                                                                                                                                                                 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 other;
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  16-APR-1999
119-APR-1999
21-APR-1999
23-APR-1999
23-APR-1999
30-APR-1999
30-APR-1999
30-APR-1999
30-APR-1999
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30-APR-1999
30-APR-1999
30-APR-1999
31-APX-1999
31-AP
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09-MAR-1999;
23-MAR-1999;
25-MAR-1999;
29-MAR-1999;
01-APR-1999;
06-APR-1999;
08-APR-1999;
99US-01123180

99US-0125788

99US-0125788

99US-012664

99US-012664

99US-0128746

99US-0130677

99US-0130691

99US-0132486

99US-0132486

99US-0134286

99US-013421

99US-013425

99US-013425

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99US-0140823

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RESULT 6
AAA79696/c
ID AAA79696 standard; cDNA; 2084 BP
XX
AC AAA79696;
XX
DT 27-NOV-2000 (first entry)
XX
Eucalyptus grandis cell signalli
XX
Eucalyptus grandis; pinus radiat
KW plant cell signalling; modulatli
XX
plant cell signalling; modulatli
KW environmental change; developmen
KW environmental change; developmen
KW elongation; survival; disease re
XX
PN W0200042171-A1.
XX
PN W0200042171-A1.
XX
PD 20-JUL-2000.
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11-JAN-1999; 99US-0162866.
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PR 01-NOV-1999; 99US-0162866.
YX
CENESIS RES & DEV CORP L
XX
PI (Strabala TJ, Nieuwenhuizen NJ;
XX
Strabala TJ, Nieuwenhuizen NJ;
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Best Local S
Matches 20
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                                                                                                                                                                                                 Eucalyptus grandis; Pinus radiata; Monters
plant cell signalling; modulation; transge
environmental change; development; cell pu
elongation; survival; disease resistance;
                                       (GENE-) GENESIS RES & DEV CORP LTD
                                                                                                                                                                                                                                                                                                                                                                                                  690
                                                                                                                                                                                                                                                                                                                                                                                               tgttggatcacaagatttgata
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity 90.9
20; Conservative
                                                                                                                                                                                                                                                              grandis cell signalling
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9908-0158262

9908-0158232

9908-0159294

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9908-0159329

9908-0159330

9908-01596331

9908-01596331

9908-0160767

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9908-0160140

9908-0160140

9908-0160140

9908-0161405

9908-0161405

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9908-0161359

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9908-0161359
                Nieuwenhuizen NJ;
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                                                                                                                                                                                                                                                                                                                                        ВР
                                                                                                                                                                                                              Monterey pine; plant; modification; transgenic plant; pathogen; growth; cell proliferation; differentiation;
                                                                                                                                                                                                                                                              involved
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                                                                                                                                                                                                     nutrient metabolism; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI;
             Claim 1; SEQ ID 22189;
                                                                                                                                                                                                                                                                                                EP1033401-A2
                                                                                                                                                                                                                                                                                                                                                              gene
                                                                                                                                                                                                                                                                                                                                                                             Human; 5' EST;
                                                                                                                                                                                                                                                                                                                                                                                                            Human secreted
                                                                                                                                                                                                                                                                                                                                                                                                                                         06-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAC18114;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAC18114 standard; cDNA; 447 BP:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 2084 BP; 502 A; 434 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1; Page 408-409; 527pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Isolated polynucleotide encoding a polypeptide involved in cell signaling used for generating transgenic plants with modified responses to external signals - \,
                                            diagnostic, forensic,
                                                         New nucleic acid that is a 5' expressed sequence tag (5' EST) for obtaining cDNAs and genomic DNAs that correspond to 5'ESTs and fo
                                                                                                                                       Dumas Milne Edwards
                                                                                                                                                                    (GEST ) GENSET
                                                                                                                                                                                                                                  21-FEB-2000;
                                                                                                                                                                                                                                                                 06-SEP-2000
                                                                                                                                                                                                                                                                                                                              Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                         therapy;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                  2000EP-0200610
                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                           chromosome mapping;
                                                                                                                                                                                                                                                                                                                                                                                                       protein 5' ESŢ,
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                                          gene therapy and chromosome
             71pp + CD-ROM; English
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Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                          SEQ ID NO: 22189:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          566 G; 582 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                              tag; secreted protein; cDNA isolation;
ss.
                                                                                                                                       Giordano
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                                          mapping procedures
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RESULT
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Best Local
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            23 APR 1999;
23 APR 1999;
28 APR 1999;
30 APR 1999;
04 APR 1999;
04 - MAY 1999;
05 - MAY 1999;
06 - MAY 1999;
07 - MAY 1999;
11 - MAY 1999;
                                                                                                                                                     23-MAR-1999;
25-MAR-1999;
29-MAR-1999;
01-APR-1999;
06-APR-1999;
16-APR-1999;
16-APR-1999;
19-APR-1999;
21-APR-1999;
                                                                                                                                                                                                                                                                        05-MAR-1999;
09-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                  metabolic
                                                                                                                                                                                                                                                                                                                                                                                                                                            Hybridisation assay; genetic mapping; gene expression control; protein identification; signal transduction pathway;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAC42365;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAC42365 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 447 BP; 187 A; 57 C; 66 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       expression and secretion vectors.
                                                                                                                                                                                                                                                                                                                                                                                                        Arabidopsis thaliana
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                                                                                                                                                                                                                                                                                                                                                                                                                               pathway; promoter;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
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99US-0127462.
99US-0128714.
99US-0128714.
99US-0129845.
99US-0130077.
99US-0130510.
99US-01304891.
99US-0131449.
99US-0132487.
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99US-0123548.
99US-0125788.
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99US-0126785
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA;
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86.4%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                  termination sequence;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 17.2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQ
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1999; 99US 1999; 99US 1999; 99US 1999; 99US	21-OCT-1999; 99US 21-OCT-1999; 99US 21-OCT-1999; 99US	21-OCT-1999; 990S	14-0CT-1999; 99US	14-0CT-1999; 99US	14-OCT-1999; 99US	13-OCT-1999; 99US	13-OCT-1999; 99US	08-OCT-1999; 99US	07-OCT-1999; 99US	05-OCT-1999; 99US	04-OCT-1999; 99US	28-SEP-1999; 99US	24-SEP-1999; 99US	23-SEP-1999; 99US	20-SEP-1999; 99US	16-SEP-1999; 99US	13-SEP-1999; 99US	10-SEP-1999; 99US	01-SEP-1999; 99US	31-AUG-1999; 99US	27-AUG-1999; 99US	27-AUG-1999; 99US 27-AUG-1999; 99US	26-AUG-1999; 99US	23-AUG-1999; 99US	23-AUG-1999; 99US	R 20-AUG-1999; 99US	R 20-AUG-1999; 99US	18-AUG-1999; 99US	16-AUG-1999; 99US	R 13-AUG-1999; 99US	R 12-AUG-1999; 99US	R 11-AUG-1999; 99US	R 10-AUG-1999; 990S	R 09-AUG-1999; 99US	06-AUG-1999; 99US	05-AUG-1999; 99US	05-AUG-1999; 99US	04-AUG-1999; 99US	03-AUG-1999; 99US	02-AUG-1999; 99US	R 02-AUG-1999; 99US	R 28-JUL-1999; 99US	R 27-JUL-1999; 99US	R 27-JUL-1999; 99US
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02-AUG-1999; 03-AUG-1999; 04-AUG-1999;

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RESULT 1
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Best Local :
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28-OCT-1999;
29-OCT-1999;
                                                                                                                                  The invention relates to polynucleotides encoding novel human proteins or their active domains. The polypeptides, polynucleotides and antibodies raised against the polypeptides are used in a method of treatment of a mammal and prevention of disorders caused by the aberrant protein expression or activity. The polypeptides can be used as molecular weight markers, food supplements, and in antibody production. The polypeptides are used to identify compounds which bind to the polypeptides. Polynucleotides of the invention are used as probes and primers, for sequencing, for chromosome or gene mapping, in the production of recombinant proteins, and in generating anti-sense DNA or RNA and in gene therapy. Polypeptides of the invention can be used to target drugs to a tumour, in assays to determine biological activity, to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; novel protein; ss; Antianaemic; osteopathic; antiinflammatory; immunomodulatory; cytostatic; neuroprotective; vulnerary; nootropic; anticonvulsant; antiarthritic; cerebroprotective; antifungal; antiviral; antibacterial; antiallergic; dermatological; haemostatic; antiasthmatic; thrombolytic; immunogen; antibody; gene therapy; neurological disorder; parkinson's disease; inflammatory disorder; cancer; asthma; osteoporosis
                             raise antibodies/elicit an immune response, to determine quantitative protein levels, as tissue markers, and to isolate receptors or ligands. Polypeptides of the invention may also be useful in treating platelet disorders, stem cell disorders, regenerating bone, cartilage, tendon, ligament and/or nerve tissue, wound healing, treating burns, promoting the proliferation, differentiation and survival of tem cells, as a
                                                                                                                                                                                                                                                                                                                                                                            Claim
                                                                                                                                                                                                                                                                                                                                                                                                             Isolated polypeptides useful for treating anti-inflammatory diseases, nervous system disorders, and for regenerating bone and cartilage -
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contraceptive, Alzheimer's, P
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86.4%;
                  osteoporosis and osteoarthritis, anaemia,
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Pred. No. 36;
Huntington's diseases, amylotrophic
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אקריקים של היא האים האלים האלי

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23-AUG-1999
23-AUG-1999
23-AUG-1999
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99US-0149929. 99US-0149902. 99US-0149930.

990S-0148565.
990S-0148684.
990S-0149175.
990S-0149175.
990S-0149426.
990S-0149722.
990S-0149723.

99US-0147935. 99US-0148171. 99US-0148319. 99US-0148341.

99US-0147416 99US-0147493 99US-0147303

9999988

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RESULT 11
AAH15352/c
ID AAH15352 standard; cDNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                          27-AUG-1999;
11-JAN-2000;
02-MAY-2000;
09-JUN-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         screrosis, stroke, immune deficiencies resulting from bacterial, fungal infection or from autoimmunity, cancer, allergy, asthma, graft-versus-host disease, eczema, haemophilia, thrombosis, anti-inflammatory diseases, nervous system disorders, and infection present sequence encodes a protein of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1929
        sequence and an oligonucleotide comprising a sequence complementary to polynucleotide which comprises a 3'-end sequence, where the oligonucleotide comprises at least 15 nucleotides and the combination the 5'-end sequence/3'-end_sequence is selected from those defined in the specification. The primer sets can be used in antisense therapy and in gene therapy. The primers are useful for synthesising polynucleotid marticulary.
                                                                                                                     full-length cDNAs defined in the specification. Where a primer set comprises: (a) an oligo-dT primer and an oligonucleotide complementary to the complementary strand of a polynucleotide which comprises one of the 5602 nucleotide sequences defined in the specification, where the oligonucleotide comprises at least 15 nucleotides; or (b) a combination of an oligonucleotide comprises at least 15 nucleotides; or (b) a combination
                                                                                                                                                                                                                                                                                                                            WPI;
                                                                                             of an oligonucleotide comprising a sequence complementary to the complementary strand of a polynucleotide which comprises a 5'-end
                                                                                                                                                                                                                            Claim
                                                                                                                                                                                                                                                                  Primer sets for synthesizing polynucleotides, particularly the 5602 full-length cDNAs defined in the specification, and for the detecti and/or diagnosis of the abnormality of the proteins encoded by the
                                                                                                                                                                                                                                                                                                                                                                                                 (HELI-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  29-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             28-JUL-2000; 2000EP-0116126
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       07-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EP1074617-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human cDNA sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAH15352;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 2038
                                                                                                                                                                                                                                                        full-length cDNAs
                                                                                                                                                                                                                                                                                                                                                        Ishii S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 tgttggatcacaagatttgata
                                                                                                                                                                                                present invention
                                                                                                                                                                                                                                                                                                                            2001-318749/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            tgttggagcacaagattttaaa 1950
                                                                                                                                                                                                                            8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19;
                                                                                                                                                                                                                                                                                                                                                                    Isogai T,
                                                                                                                                                                                                                                                                                                                                                                                                HELIX RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                            SEQ
                                                                                                                                                                                                                                                                                                                                                    sogai T, Nishikawa
Sugiyama T, Wakama
                                                                                                                                                                                                                                                                                                                                                                                                                          99JP-0300253.
2000JP-0118776.
2000JP-0183767.
2000JP-0241899.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                          ID 13526;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        detection; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 99JP-0248036.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       658
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ ID NO:13526
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              78.2%;
86.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Α,
                                                                                                                                                                                                describes primer sets
                                                                                                                                                                                                                                                                                                                                                       Wakamatsu
                                                                                                                                                                                                                         2537pp +
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       400 C;
 CDNAS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pred. No. 38;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                      ₽,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ВP
                                                                                                                                                                                                                            G
The primers are
                                                                                                                                                                                                                                                                                                                                                                                                                                      ...
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                                                                                                                                                                                                                                                                                                                                                     Hayashi K,
A, Nagai K,
                                                                                                                                                                                                                            ROM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      antisense therapy; gene therapy; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       477
                                                                                                                                                                                                                          English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       for synthesising 5602
                                                                                                                                                                                                                                                                                                                                                     Saito K,
Otsuki
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0 other;
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                                                                                                                                                                                                                                                                                                                                                        Ŧ
                                                                                                                                                                                                                                                                                                                                                                      Yamamoto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2038;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               infection
                                                                                                                                                                                                                                                                  detection by the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
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AAT90543/c
ID AAT90543
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       detection and/or diagnosis of the abnormality of the proteins encoded by the full-length cDNAs. The primers allow obtaining of the full-length cDNAs easily without any specialised methods. AAH03166 to AAH13628 and AAH13633 to AAH18742 represent human cDNA sequences; AAB92446 to AAB95893 represent human amino acid sequences; and AAH13629 to AAH13632 represent oligonucleotides, all of which are used in the exemplification of the present invention.
This DNA sequence comprises the maize Z33 ZM41A gene. This is an orthologue of Arabidopsis Ms41-A (see AAT90522), a gene that confers male fertility. It was obtained following a database search for sequences that showed homology to Ms41-A DNA, isolation of a partial cDNA clone, and use of this clone to isolate Zm41-A genes Z31 (AAT90542), Z33 (AAT90543) and Z35 (AAT90544) from genomic lambda libraries. The genes show a high level of conservation. Z35 may
                                                                                                        Nucleic acid encoding proteins - used to control fertility and
                                                                                                                                                                                                                                                                                                          exon
                                                                                                                                                                                                                                                                                                                                                    exon
                                                                                                                                                                                                                                                                                                                                                                                                        Key
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAT90543;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                         WPI; 1997-351055/32.
                                                                                                                                                                                                                                20-DEC-1996;
                                                                                                                                                                                                                                                      03-JUL-1997
                                                                                                                                                                                                                                                                          W09723618-A1
                                                                                                                                                                                                                                                                                                                                                                       intron
                                                                                                                                                                                                                                                                                                                                                                                                                             Zea mays
                                                                                                                                                                                                                                                                                                                                                                                                                                                  ribozyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             2m41-A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Maize male
                                                                                                                                                                                     (GENE-) GENE SHEARS PTY LTD.
                                                                                                                                                                                                                                                                                                                              intron
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      783 TGTTGGATGACCAGATTTGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 tgttggatcacaagatttga
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                l Similarity
18; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                             male
                                                                                     Fig
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1764
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 male sterile; maize; Ms41-A; ds.
                                                                                                                                                              Garcia D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               fertility gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                   15; 85pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP;
                                                                                                                                                                                                                               96WO-GB03191
                                                                                                                                                                                                                                                                                                                           /*tag= b
936..1090
/*tag= c
1091..1167
                                                                                                                                                                                                                                                                                                                                                                      /*tag= a
916..935
                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers 823..915
                                                                                                                                                                                                                                                                                                          1168.
                                                                                                                                                                                                                                                                                                                    /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     511 A; 347 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DNA;
                                                                                                                                                                                                                                                                                                         g= d
..1332
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        76.48;
90.08;
                                                                                                                                                               Hodge
                                                                                     English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        764
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Z33 Zm41-A
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                                                                                                                                                            ₽,
                                                                                                          and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score
                                                                                                                   involved
                                                                                                                                                                                                                                                                                                                                                                                                                                                           seed; transgenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      392
                                                                                                          for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                              Perez
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     G;
                                                                                                        lved in male fertility in plants production of hybrid seed
                                                                                                                                                               ٦,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            58;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DВ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            plant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
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Query Match
Best Local Similarity
Watches 18; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           be derived from 231 via genetic rearrangements, deletions and/or insertions. 233 has subsequent deletions from 235 and is truncated, having only exons 3, 5 and 6. The Ms41-A and Zm41-A genes, antisense or ribozyme sequences can be used to produce transgenic plants with controlled male fertility. Male sterile plants are useful for hybrid seed production, particularly in plants where restoration of fertility is not needed, e.g.
                     Eucalyptus grandis or Pinus radiata. The present sequence is the coding sequence for one such transcription factor. The transcription factor may be used to produce a plant having modified gene expression such as a woody plant e.g. a eucalyptus, pine, acacia, poplar, sweetgum, teak, or mahogany species or to modify the activity of a polypeptide in a plant. The transcription factors of the present invention are members from the following families of regulatory proteins: bZIP, bZIP family of G-box binding factors, basic helix-loop-helix zipper.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Plant; transcription factor; gene expression; eucalyptus; pine; poplar; sweetgum; teak; mahogany; bZIP; G-box binding factor; basic helix-loop-helix zipper; homeotic; homeodomain; homeobox; homeodomain zipper; LIM domain; AP2; EREBS; zinc finger domain; type 2 Cys2His2; CCAAT box element; MYB: ss
                                                                                                                                                                             Claim
                                                                                                                                                                                                                    New isolated polynucleotide encoding a plant transcription factor for producing a plant e.g. a woody plant, preferably eucalyptus or pine, having modified gene expression or modified activity of a polypeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAC56059
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Brassicaceae,
 and
                                                                                                                                             The present invention relates to novel plant transcription factors
                                                                                                                                                                                                                                                                                WPI;
                                                                                                                                                                                                                                                                                                           Wood M,
                                                                                                                                                                                                                                                                                                                              (GENE-) GENESIS RES & DEV CORP LTD (FLET-) FLETCHER CHALLENGS FORESTS
                                                                                                                                                                                                                                                                                                                                                                                                                                09-MAR-2000; 2000WO-US06112
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200053724-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eucalyptus grandis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eucalyptus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    25-JAN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAC56059 standard;
                                                                                                                                                                                                                                                                                                                                                                                    11-MAR-1999;
18-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              14-SEP-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    268
             nomeotic/homeodomain/homeobox/MADS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 tgttggatcacaagatttga 20
 EREBS,
                                                                                                                                                                                                                                                                              2000-579369/54
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TGTTGAATCACAAGATCTGA 249
                                                                                                                                                                            1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2582
                                                                                                                                                                            Pages
                                                                                                                                                                                                                                                                                                          McGrath A,
 zinc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   B₽;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               lettuce, spinach and onions.
                                                                                                                                                                          94-95; 747pp; English
                                                                                                                                                                                                                                                                                                                                                                                  99US-0266513.
99US-0149485.
 finger
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   670 A; 512
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       transcription
                                                                                                                                                                                                                                                                                                           Shenk MA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         76.4%;
90.0%;
 domains
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2444
                                                                                                                                                                                                                                                                                                                                       FORESTS LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score
Pred.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP
of type 2 Cys2His2,
                                                                                                                                                                                                                                                                                                           Glenn M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   537
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     factor DNA sequence #190.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16.8;
No. 60;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ç;
homeodomain zipper, LIM domain, A ype 2 Cys2His2, CCAAT box elements
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   863 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2582;
               domain, AP2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MADS;
                                                                                                                                             from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
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RESULT 14
ADA 33702/c
ID AN 4337
XX AN 4337
XX AN 4337
XX Human;
XX Human;
XX Homo s
XX WO2001
XX 29-SEI
XX 29-SEI
XX 29-SEI
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Best Local
                                                                                                                                                    expression. For example, N and P may be used to treat disorders associated with decreased expression by rectifying mutations or deletions in a patient's genome that affect the activity of P by expressing inactive proteins or to supplement the patients own production of P. Additionally, N may be used to produce the colon cancer-associated Ps. by inserting the nucleic acids into a host cell and culturing the cell to express the proteins. N and P can be used in the prevention, diagnosis and treatment of colorectal carcinomas and cancers. AAH37196 to AAH37204 and AAB77789 represent sequences used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1451
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                                                                                                                                                                                                                                                                                                                                                                                           the proteins are collectively known as colon cancer antigens. The colon cancer antigens have cytostatic activity and can be used in gene therapy and vaccine production. N and P may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                       present invention.

N.B. Pages 666 to 682 and page 7053 of the missing at time of publication, meaning no SEQ ID NO:1027 to 1052, 7921 and 7922.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAH32943 to AAH37195 and AAG73514 to AAG77788 represent human cancer-associated nucleic acid molecules (N) and proteins (P).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 1; Page 2738; 9803pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nucleic acids encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           P-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ruben
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           29-SEP-1999;
03-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28-SEP-2000; 2000WO-US26524
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 colorectal carcinoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human colon cancer antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                03-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAH33702;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (HUMA-) HUMAN GENOME SCI INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          colon cancer; colon cancer antigen; diagnosis; detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             l Similarity
17; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           for
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        preventing,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Barash SC,
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99US-0163280.
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94.4%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4277 human colon cancer-associated polypeptides, diagnosing and/or treating colorectal cancers -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Birse CE,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           G;
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                                                                                  sequence listing were 
sequences are present
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Best Local S
Matches 18
06-JUN 1997

05-SEP 1997

05-JUN 1997

06-JUN 1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; secreted protein; fusion protein; gene therapy; protein therapy; diagnosis; tissue; cancer; tumour; neurodegenerative disorder; leukaemia; developmental abnormality; foetal deficiency; blood; allergy; renal; ds; immune system; asthma; lymphocytic disease; brain; hepatic; lymphoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                   18-DEC-1997
06-JUN-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         04-JUN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          endocrine; metabolism; regulation; malabsorption; gastritis; neoplasm
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cognitive disorder; schizophrenia; prostate; obesity; osteoclast; thymus; osteoporosis; arthritis; testis; lung; thyroiditis; thyroid; digestion;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human secreted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-MAR-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAV84598 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             immune system; asthma; lymphocytic disease; brain; hepatic; lymphoma; inflammation; ischaemic shock; Alzheimer's disease; restenosis; AIDS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           122 TGTTGTAGCACAAGATTTTAT 102
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970S-00488949
970S-0048915
970S-0048949
970S-0048972
970S-0049375
970S-0049375
970S-0057644
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970S-0048878
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97US-0048895.
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97US-0048915.
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97US-0048881.
97US-0048884.
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85.7%;
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Pred. No. 96;
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05-SEP-1997
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                                                                                                                                                                                                                     (HUMA-) HUMAN
                                                                                                                                                                                                         Carter KC,
                                                                                                                                                                                                                      GENOME SCI INC.
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97US-00488876.
97US-0048880.
97US-0048883.
97US-0048893.
97US-0048891.
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97US-0057662.
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97US-0057654.
97US-0057656.
97US-0057666.
97US-0057760.
97US-0057769.
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Fan P, Florence K, Li Y, Moore Shi Y, Sopp 1999-059865/05. DB; AAW88721. Feng P, Ferrie AM, Fischer ce K, Greene JM, Hu J, Kyz Moore PA, Ni J, Olsen HS, Soppet DR, Wei Y, Young P, llon PJ, Ebner R, Enu..., Fischer CL, Florence C;
T Kyaw H, Lafleur DW;
Kyaw H, CA, Ruben SM; Young P, Rosen CA, R

New isolated human genes and the secreted polypeptides they encode useful for diagnosis and treatment of e.g. cancers, neurological disorders, immune diseases, inflammation or blood disorders

Claim Page 450-451; 772pp; English.

The invention relates to nucleic acid sequences (AAV84411 to AAV84633) CC encoding human secreted proteins (AAW88334 to AAW88756). The secreted protein gene sequences are deposited with the ATCC under deposit numbers CC ATCC 97979, 97974, 97975, 97976, 97977, 209007, 209008, 209009, 209010, CC 209011, 209008, 2090081, 209002, 209008, 2090010, CC 209011, 209008, 2090081, 209008, 2090010, CC 209011, 209008, 2090081, 209008, 2090010, CC 209011, 209008, 2090081, 209008, 2090010, CC 209011, 2090081, 2090081, 2090082, 2090083, 2090084, 2090085, 209511. HOST CC 209011, 2090081, 2090081, 2090082, 2090083, 2090083, 2090085, 209511. HOST CC 209011, 2090081, 2090081, 2090082, 2090081, 2090010, CC 209011 secured and amino acid sequences are useful for are CC useful for gene therapy, retaining a medical conditions e.g. by protein or gene therapy, Pathological conditions can be also CC diagnosed by determining the amount of the new polypeptides in a sample CC or by determining the presence of mutations in the new polynucleotides. Specific uses are described for each of the polynucleotides, based on CC which tissues they are most highly expressed in, and include developing products for the diagnosis or treatment of cancer, neurodegenerative CC disorders, tumours, leukemias, diseases of the immune system, autoimmune disorders, hepatic and renal disease, lumphomas, inflammation, allergies, to the complete and the content of the complete and to the content of the complete and to the content of the complete and the content of the content of the content of the complete and the content of the ischemic shock, cognitive disorders,

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CC restenosis, prostate diseases, obesity, disorders involving osteoclasts
CC such as osteoporosis, arthilis or malignancies, diseases of testes,
CC lung or thymus, digestive/endocrine disorders, infections and AIDS. The
CC polypeptides are also useful for identifying their binding partners.
CC The present sequence represents a gene encoding a human secreted protein
CC (see descriptor line for gene number and clone identification).

XX
SQ Sequence 951 BP; 296 A; 134 C; 192 G; 329 T; 0 other;

Query Match
Best Local Similarity 85.7%; Pred. No. 1.1e+02;
Matches 18; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

QU 1 tgttggatcacaagatttgat 21
QU 1 tgttggatcacaagatttgat 21
QU 1 tgttggatcacaagatttgat 21
Db 524 TGTTGGTTCTCAAGATCTGAT 504

Search completed: December 15, 2001, 03:29:43
Job time: 7883 sec
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US-08-014-326-1
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; Sequence 31, Application

; Patent No. 6207883

; GENERAL INFORMATION:
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6 5256642-3 6 5472939-3 7 Sequence 6 5256642-1 6 5472939-1 6 5472939-1 7 Sequence 7 Sequence 8 525642-1 8 Sequence 9 Sequence 9 Sequence 10 Sequence	66.4	66.4	67.3	67.3	67.3	67.3	67.3	67.3	67.3	67.3	67.3	67.3	67.3	69.1	69.1	69.1	69.1	69.1
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	Sequence 17, Appl	13,	! 1, A	28,	9	Sequence 5, Appli	Sequence 9, Appli	Sequence 3, Appli	Sequence 3, Appli	Sequence 3, Appli	4, 1	71,	13,	•	•		No.	NO.

ALIGNMENTS

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US-08-014-326-1
Sequence 1, Application US/08014326
Patent No. 549854
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 90.0
Matches 18; Conservative
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APPLICATION NUMBER: WO
FILING DATE:
INFORMATION FOR SEQ ID NO:
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LENGTH: 2582 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
               APPLICANT: Gengenbach, I
APPLICANT: Somers, David
APPLICANT: Wyse, Donald
APPLICANT: Gronwald, Jol
APPLICANT: Egli, Margare
APPLICANT: Lutz, Shiela
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOTTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compati
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 TITLE OF INVENTION:
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             Wyse, Donald L. Gronwald, John W. Egli, Margaret A. Lutz, Shiela M.
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Somers, David A.
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Patent No. 6069298
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Best Local Similarity 85.7%;
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APPLICATION NUMBER: US 07/917,462
FILING DATE: 21-JUL-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/538,674
FILING DATE: 18-JUN-1990
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/269,584
FILING DATE: 10-NOV-1988
ATTORNEY,AGENT INFORMATION:
                                                                                                                                                                                                                                       GENERAL INFO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: 612-332-9081
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                   TITLE OF INVENTION: METHODS AND AN ACETYL COA CARBOXYL.
TITLE OF INVENTION: FOR CONFERRING HERBICIDE TOLERANCE
TITLE OF INVENTION: OIL CONTENT OF PLANTS
NUMBER OF SEQUENCES: 11
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30 (EPO).
CURRENT APPLICATION DATA:
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION: TELEPHONE: 612-332-5300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0.
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CITY: Minneapolis
STATE: MN
                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Kowalchyk, Katherine M. REGISTRATION NUMBER: 36,848 REFERENCE/DOCKET NUMBER: 600.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY:
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APPLICATION NUMBER: U
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                                                                                                                                                                                                                                                            INFORMATION:
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00 No. 5498544west Center
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05-FEB-1993
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                                                                                                                                                                         METHODS AND AN ACETYL COA CARBOXYLASE GENE
FOR CONFERRING HERBICIDE TOLERANCE AND AN ALTERATION IN
OIL CONTENT OF PLANTS
    US/08/417,089
D: 1:
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Pred. No. 2
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; MOLECULE TYPE:
US-08-417-089-1
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Best Local
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                                                                                                                                                                                                CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UNMBER: 08/417089
FILING DATE: 05-APR-1995
APPLICATION NUMBER: 08/014326
FILING DATE: 05-FEB-1993
APPLICATION NUMBER: 07/917462
FILING DATE: 21-UUL-1992
APPLICATION NUMBER: 07/538674
FILING DATE: 18-UUN-1990
ATTORNEY/AGENT INFORMATION:
AMMER WOSSGROF, WATTEN D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                              INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 2001 base pairs
TYPE: nucleic acid
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OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/695,
                                                                                                                       REGISTRATION NUMBER: 30,440
REFERENCE/DOCKET NUMBER: 600
TELECOMMUNICATION INFORMATION:
TELEPHONE: 612-373-6900
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                                                             SEQUENCE CHARACTERISTICS:
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STRANDEDNESS: single
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              STRANDEDNESS:
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TOPOLOGY:
                                                                                                             TELEFAX: 612-339-3061
                                              LENGTH:
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18; Conservative
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                                              2001 base pairs
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Wyse, D.
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Lutz, S. M
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                single
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Pred. No. 27;
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Best Local S
Matches 18
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                                                                                                                                                         Matches
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FILING DATE: 13-APR-1998
CLASSIFICATION: 800
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US96/04625
FILING DATE: 04-APR-1996
ATTORNET/AGENT INFORMATION:
NAME: Embretson, Janet E.
REGISTRATION NUMBER: 39,665
REGISTRATION NUMBER: 39,665
                                                                                                                                                                                                                                                                                                                                                                                                       REFERENCE/DOCKET NUMBER: 39,665
TELECOMMUNICATION INFORMATION:
TELEPHONE: 612-339-0331
TELEFAX: 612-339-3061
TELEFAX: 612-339-3061
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                                                                                                                                                                                                                                                                      TOPOLOGY: linear MOLECULE TYPE: cDN HYPOTHETICAL: NO ANTI-SENSE: NO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEO Version 1.5
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                            FRAGMENT TYPE:
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 2001 base pairs
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CITY: Minneapolis
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                                                                                                                                                                    Local Similarity
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85.7%;
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Pred. No. 27;
                                                                                                                                                                       Score 16.2;
Pred. No. 27;
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ACETYL COA CARBOXYLASE GE
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; MOLECULE TYPE:
US-08-695-421-1
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                                                                                                                                                                                                                                      В
                                                                                                                                                                              RESULT
             Sequence 1, Application US/08244537
Patent No. 5854420
GENERAL INFORMATION:
APPLICANT: ASHTON, ANTHONY R.
APPLICANT: JENKINS, COLLN L.D.
APPLICANT: WHITFELD, PAUL R.D.
APPLICANT: WHITFELD, PAUL R.D.
TITLE OF INVENTION: MAIZE ACETYL COA CARBOXYLASE ENCODING
TITLE OF INVENTION: DNA CLONES
                                                                                                                                                                                                                                                                                                         Query Match 73.6%;
Best Local Similarity 85.7%;
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION: 800

PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/417089

FILING DATE: 05-APR-1995

APPLICATION NUMBER: 08/014326

FILING DATE: 05-FEB-1993

APPLICATION NUMBER: 07/917462

FILING DATE: 21-UU-1992

APPLICATION NUMBER: 07/538674

FILING DATE: 18-UN-1990

ATTORNEY/AGENT INFORMATION:
NAME: WOESSNEY, WATTON DATE: 18-UN-1990

ATTORNEY/AGENT INFORMATION:
NAME: WOESSNEY, WATTEN D

REFIERENCE/DOCKET NUMBER: 600.311

TELECOMMUNICATION INFORMATION:
TELEPHONE: 612-373-6900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO:
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APPLICANT:
APPLICANT:
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SOFTWARE: FastSEQ Version 2.0
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Diskett
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: METHODS AND AN TITLE OF INVENTION: FOR CONFERRING
NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: nucleic acid
STRANDEDNESS: sing
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ZIP: 55402
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH:
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Somers, D. A.
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linear
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IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Schwegman,
.O. Box 2938
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           23-AUG-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     600.318US2
                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                           Score 16.2;
Pred. No. 2
                                                                                                                                                                                                                                                                                                          Mismatches
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HERBICIDE TOLERANCE AND
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) AN ALTERATION IN OIL
                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                          Gaps
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ADDRESSEE: STREET: 11 CITY: Wash STATE: D.

Washington

E: CUSHMAN DARBY & CUSHMAN, L.L.P. 1100 New York Avenue, N.W.

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US-08-417-089-5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 73.6%; Score 16.2;
Best Local Similarity 85.7%; Pred. No. 31
Matches 18; Conservative 0; Mismatch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION: 800
PRIOR APPLICATION DATA:
APPLICATION UNMBER: GB 9125330.2
FILING DATE: 28-NOV-1991
PRIOR APPLICATION DATA:
APPLICATION UNMBER: PCT/GB92/02205
FILING DATE: 27-NOV-1992
ATFORNEY/AGENT INFORMATION:
NAME: KOKULIS, PAUL N.
REGISTRATION NUMBER: 16,773
REFERENCE/DOCKET NUMBER: 203094/SE
                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                             APPLICATION NUMBER: UINFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEPHONE: (202) 861-3000
TELEFAX: (202) 822-0944
TELEX: 6714627 CUSH
INFORMATION FOR SEQ ID NO: 1:
TOPOLOGY: 1i
MOLECULE TYPE:
8-417-089-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                  SOFTWARE: PatentIn Rel CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                               TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                           COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                            SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/0 FILING DATE: 18-AUG-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: U.S.A. ZIP: 20005-3918
                                                                                                                                                                                                  COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY:
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                                                  STRANDEDNESS:
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                                                                                        ENGTH:
                                                               nucleic acid
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                 DNA (genomic)
                                                    single
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FOR CONFERRING HERBICIDE TOLERANCE AND AN ALTERATION IN
OIL CONTENT OF PLANTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PCT/GB92/02205
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Mismatches
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Patent No. 6146867
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local
                                               Query Match 73.6
Best Local Similarity 85.7
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM Compatible
COMPUTER: DOS
COMPANTING SYSTEM: DOS
SOFTWARE: FASTSEQ Version 2.0
SOFTWARE: FASTSEQ Version 2.0
CURRENT APPLICATION NUMBER: US/08/695,65
FILING DATE: 12-AUG-1996
CLASSIFICATION: 435
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/417089
FILING DATE: 05-APR-1995
APPLICATION NUMBER: 08/014326
FILING DATE: 05-FBB-1993
APPLICATION NUMBER: 07/917462
FILING DATE: 21-JUL-1992
APPLICATION NUMBER: 07/538674
FILING DATE: 18-JUN-1990
ATTORNEY/AGENT INFORMATION
NAME: WOESSNEY, WAITEN D
REGISTRATION NUMBER: 30,440
PRESISTRATION NUMBER: 30,440
PRESISTRATION NUMBER: 30,440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ZIP: 55402
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
ADDRESSEE: Schwegman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Egli, M. APPLICANT: Lutz, S. TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3972 GTTGGATGACAAGAGTTGTTA 3992
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                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                               TELEPHONE: 612-373-6900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
                                                                                                                                                MOLECULE TYPE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: P.O. Box CITY: Minneapolis
                                                                                                                                                                           STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY:
                                                                                                                                                                 TOPOLOGY:
                                                                                                                                                                                                                                                                              TELEFAX: 612-339-3061
                                                                                                                                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE: Diskette
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2 gttggatcacaagatttgata
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                                                                                                                                                                                                           7470 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gronwald,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gengenbach,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Wyse, D. L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Somers, D. A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                 linear
                                                                                                                                                  CDNA
                                                                                                                                                                             single
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                                                              73.6%;
85.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            В. G.
                  22
                                               Score 16.2; D
Pred. No. 32;
0; Mismatches
                                                                                                                                                                                                                                                                                                                               600.318US3
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Pred. No. 32;
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                                                                                Length 7470;
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US-08-930-285-5 ; Sequence 5, Application US/08930285 ; Patent No. 6222099 ; Patent No. 6222099

GENERAL INFORMATION:

CORRESPONDENCE ADDRESS:

CITY: Minneapolis STATE: MN

STREET:

ADDRESSEE:

SE: Schwegman, Lundberg, Woessner & Kluth, P. A. P. O. Box 2938

COUNTRY:

USA

55402 ĭ APPLICANT: Regents of the University of Minnesota, et al.
TITLE OF INVENTION: TRANSGENIC PLANTS EXPRESSING ACETYL COA CARBOXYLASE GE
NUMBER OF SEQUENCES: 26

В

3972 GTTGGATGACAAGAGTTGTTA 3992

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                                                                                                                    US-08-695-421-5
                                                                                                                               RESULT
                                                                                             Sequence 5,
Patent No. 6
                                                                                                                                                                                                                  Matches 18;
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Best Local Similarity
                                                                                 GENERAL INFORMATION:
                                                                                                                                                                    3972 GTTGGATGACAAGAGTTGTTA 3992
                                                            APPLICANT:
                                                                      APPLICANT:
TITLE OF INVENTION:
             APPLICANT:
                         APPLICANT:
                                    APPLICANT:
                                               APPLICANT:
                                                                                                                                                                                2 gttggatcacaagatttgata 22
                                                                                                         Application US/08695421
                                   Wyse, D. L.
Gronwald, J
                                                                                                                                                                                                                  Conservative
                     Egli, M. A.
                                                         Somers, D. A.
                                                                    Gengenbach,
                                                                                                                                                                                                                           73.6%;
METHODS AND AN ACETYL COA CARBOXYLASE GENE
                                                                      B. .G.
                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                            Score 16.2;
Pred. No. 32;
                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                       DB
                                                                                                                                                                                                                                      Length 7470;
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; ORIGINAL SOURCE: US-08-930-285-5

FRAGMENT TYPE:

MOLECULE TYPE: CI HYPOTHETICAL: NO ANTI-SENSE: NO

TOPOLOGY:

linear

CDNA

INFORMATION FOR SEQ ID NO:

5.

TELEFAX: 612-339-3061

SEQUENCE CHARACTERISTICS:
LENGTH: 7470 base pairs
TYPE: nucleic acid
STRANDEDNESS: single

NAME: Embretson, Janet 5.
REGISTION NUMBER: 39,665
REFERENCE/DOCKET NUMBER: 600
TELECOMMUNICATION INFORMATION:
TELECHONE: 612-339-0331

600.318US4

PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US96/04625
APPLING DATE: 04-APR-1996
ATTORNEY/AGENT INFORMATION:

OPERATING SYSTEM: DOS SOFTWARE: FastSEQ Version 1.5 CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/930,285 FILING DATE: 13-APR-1998

CLASSIFICATION: 800

COMPUTER READABLE FORM: MEDIUM TYPE: Diskett

COMPUTER:

IBM Compatible Diskette

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US-08-602-145-14/c
Sequence 14, Application US/08602145; Patent No. 602536; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                    B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; MOLECULE TYPE: CDNA US-08-695-421-5
                                                                                                                                                                                                                                                                                                                                             RESULT 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OPERATING STEEM: DOS
SOFTWARE: FastSEQ Version 2.0
CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/08/695,4:
FILING DATE: 23-AUG-1996
CLASSIFICATION ENDATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION UNMBER: 08/417089
FILING DATE: 05-APR-1995
FILING DATE: 05-APR-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                  APPLICANT: Goltry, Kristin L.
APPLICANT: Greenberger, Joel S.
TITLE OF INVENTION: DETERMINING EXPOSURE TO IONIZING RADIATION
TITLE OF INVENTION: AGENT WITH PERSISTENT BIOLOGICAL MARKERS
NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                    3972 GTTGGATGACAAGAGTTGTTA 3992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 18-JUN-1990
ATTORNEY/AGENT INFORMATION:
NAME: Woessner, Warren D
REGISTRATION NUMBER: 30,4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: FO NUMBER OF SEQUENCES: 1 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELECOMMUNICATION INFORMATION: TELEPHONE: 612-373-6900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: 08/014326
FILING DATE: 05-FEB-1993
APPLICATION NUMBER: 07/917462
FILING DATE: 21-JUL-1992
APPLICATION NUMBER: 07/538674
                                                                                    STREET: 3000 K SO
CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: UZIP: 55402
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STREET: P
                                       COUNTRY: USA
ZIP: 20007-5109
                                                                           STATE: D.C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: 612-339-306
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY:
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                                                                                                                                  ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 7470 base pairs
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P.O. Box 2938
                                                                                                            E: Foley & Lardner
3000 K Street, N.W., Suite 500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                linear
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Pred. No. 3;
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SOFTWARE: Patentin Release #1:0, CURRENT APPLICATION DATA:

APPLICATION NUMBER:

COMPUTER: IBM PC
OPERATING SYSTEM:
SOFTWARE: Patent

IBM PC compatible

PC-DOS/MS-DOS

Version #1.30

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Вb
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Best Local S
Matches 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
                                                                         FILING DATE: 13-SEP-1996
ATTORNEY/AGENT INFORMATION:
NAME: MANDIAGENT INFORMATION:
NAME: MANDIAGENT ANY E.
REGISTRATION NUMBER: 36,207
REFERENCE/DOCKET NUMBER: DFN-
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
                                            TELEFAX: (617)742-4214
INFORMATION FOR SEQ ID NO: 3:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Bent, Stephen A.
REGISTATION NUMBER: 9,768
REFERENCE/DOCKET NUMBER: 076333/0112
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)672-5300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 296 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: 1i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 15-FEB-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy of COMPUTER: IBM PC com
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Robert W. Finberg, Jeffrey M. Bergelson, APPLICANT: and Marshall S., Horwitz TITLE OF INVENTION: CAR, *A No.'6210921el Coxsackievirus and Adenovirus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
                              SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                            PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   288 TTTTGCATCATAACATTTGATA 267
                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
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TELEX: 904136
TYPE: n
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                                                                                                                                                                                         APPLICATION NUMBER:
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1095 base pairs
cleic acid
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15-FEB-1996
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Pred. No. 4
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Query Match
Best Local Similarity
Matches 18; Conserv

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0; Mismatches Score 15.6; Pred. No. 49;

70.9%;

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Length 1096;

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Matches 18; Conserv
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                          FILING DATE: 19-JUL-19 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 5.25 inch, 360 Kb storage
COMPUTER: IBM AT-compatible, 80286 processor
OPERATING SYSTEM: MS-DOS version 5.0
SOFTWARE: WordPerfect version 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FEATURE:
                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/966,040
FILING DATE: 30-DEC-1992
APPLICATION NUMBER: PCT/EP91/01361
FILING DATE: 19-JUL-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: Keil & Weinkauf
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE: CDNA
                                                                            MOLECULE TYPE:
ORIGINAL SOURCE:
                                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 1096 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
                                              FEATURE:
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CITY: V
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                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION: 435
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LOCATION:
OTHER INFORMATION: the coding region shown in (2)(ix)(B) OTHER INFORMATION: codes for the protein of SEQ ID NO: 3
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                                                               ORGANISM:
                                                                                                              TOPOLOGY:
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                                 LOCATION: 144 to 841
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Bach, Alfred
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No. 5759541el Proteins, the Preparation and Use
Thereof
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Pred. No. 49
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/928,383B
FILING DATE: 12-SEP-1997
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 60/026,100
FILING DATE: 13-SEP-1996
ATTORNEY/AGENT INFORMATION:
NAME: Mandragouras, Amy E.
REGISTRATION NUMBER: DFN-020
TELEPHONE: (617)227-7400
TELEPHONE: (617)227-7400
TELEPHONE: (617)227-7400
TELEPHONE: G17)227-7400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 15
US-08-28-383B-1/c
; Sequence 1, Application US/08928383B
; Patent No. 6210921
; GENERAL INFORMATION:
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Search completed: December 15, 2001, 03:22:46 Job time: 7561 sec
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; LOCATION: 60..1157
US-08-928-383B-1
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Best Local Similarity
Matches 18; Conserv
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MEDIIM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Robert W. Finberg, Jeffrey M. Bergelson,
APPLICANT: and Marshall S. Horwitz
TITLE OF INVENTION: CAR, A NO. 6210921el Coxsackievirus and Ademovirus
TITLE OF INVENTION: Receptor
NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS: LAHIVE & COCKFIELD, LLP
STREET: 28 State Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 tgttggatcacaagatttgata 22
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112 TGTAGTATCACAAGGTTTGCTA 91
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CITY: Boston
STATE: Massac
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ZIP: 02109
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81.8%; Pred. No. 51;
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Maximum DB seq length: 2000000000
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Perfect score:
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Maximum Match 100%

Listing first 45 summaries
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53.226 Million cell updates/sec
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Gapop 10.0 , Gapext 1.0
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Copyright (c) 1993 - 2000 Compugen Ltd.
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em_esthum:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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Result No.	Score	Query Match	Length DB	ВВ	ID	Description
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c 2	18.8	85.5	394	11	Z26559	Z26559 AT
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c 4	18.8	85.5	527	10	AV520789	AV520789 AV520789
c 5	18.8	85.5	530	10	AI995637	
ი 6	18.4	83.6	808	11	BE977618	_
c 7	18.4	83.6	934	13	CNS02BWA	
	17.8	80.9	600	1	BF006444	
ი 9	17.8	80.9	633	11	BF006254	BF006254
c 10	17.4	79.1	286	10	BB288727	BB288727 BB288727
c 11	17.4	79.1	407	10	BE581720	
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10 11 11	11 10 10	13 13 13	10 13	13 10 10	13 10 10	13 13 13		
BE118999 T20362 BI398487	C70416 AU200312 AV313332	AZ193715 CNS07DUS CNS079F4 CNS029DN CNS04HNP	AW963686 BF346432 B57784	B67515 A1777095 BE187570	AQ223477 AW034253 AI779714	W43212 AQ796989 AI733664 B62582	BE068225 AI049911 AQ595814 AQ221715	AW695022 BG448853 BE283043 BE642809 CNS00DZ6
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()		AZ193715 SP_1023_B AL440954 T7 end of AL435206 T7 end of AL187124 Tetraodon AL291166 Tetraodon	_	B67515 T22M10TR TA A1777095 EST258060 BE187570 EST336131	AQ223477 HS_2003_B AW034253 EST277824 AI779714 EST260593	W43212 22591 Lambd AQ796989 nbxb0071H AI733664 an30h03.x B62582 T22F18TR TA	BE068225 MR4-BT036 AI049911 an30h03.x AQ595814 HS_2132_B AQ221715 HS_2010_A	

ALIGNMENTS

	source	FEATURES			COMMENT	MEDLINE	JOURNAL			TITLE	AUTHORS	REFERENCE				ORGANISM	SOURCE	KEYWORDS	VERSION	ACCESSION	DEFINITION	LOCUS	RESULT 1 AV532040/c
/organism="Arabidopsis thaliana" /strain="Columbia" /db_xref="taxon:3702" /clone="FB34f09F" /clone_lib="Arabidopsis thaliana flower buds Columbia" /tissue_type="flower buds" /note="Vector: pBluescriptII SK-; Site_1: EcoRI; Site_2: XhoI"	1361	<pre>Email: asamizu@kazusa.or.jp, URL:http://www.kazusa.or.jp/en/plant/.</pre>	Kazusa DNA Research Institute Yana 1532-3, Kisarazu, Chiba 292-0812, Japan	The First Laboratory for Plant Gene Research	Contact: Erika Asamizu	20363093	DNA Res. 7, 175-180 (2000)	size-selected cDNA libraries	of 12,028 non-redundant expressed sequence tags from normalized and	A large scale analysis of cDNA in Arabidopsis thaliana: Generation	Asamizu, E., Nakamura, Y., Sato, S. and Tabata, S.	1 (bases 1 to 361)	Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.	Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;	Eukarvota: Viridiplantae: Streptophyta: Embryophyta: Tracheophyta:	Arabidopsis thaliana	thale cress.	EST.	AV532040.1 GI:8692323	AV532040	Arabidopsis thaliana		

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/tissue_type="whole seedlings"
/dev_stage="in vitro-grown etiolated seedlings,5
/note="Vector: pBluescript"
80 c 63 g 133 t 1 others
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/strain="ecotype Columbia"
/db_xref="taxon:3702"
/clone="YCVDH08"
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Kazusa DNA Research Institute
Yana 1532-3, Kisarazu, Chiba 292-0812, Japan
Email: asamizu@kazusa.or.jp, URD:http://www.kazusa.or.jp/en/plant/.
Location/Qualifiers
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A large scale analysis of cDNs in Arabidopsis thaliana: Generation of 12,028 non-redundant expressed sequence tags from normalized an size-selected cDNA libraries
                                                                                                                                                                                                    Contact: Erika Asamizu
The First Laboratory for Plant Gene Research
Kazusa DNA Research Institute
Yana 1532-3, Kisarazu, Chiba 292-0812, Japan
Email: asamizu@kazusa.or.jp, URL:http://www.kazusa.or.jp/en/plant/.
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A large scale analysis of cDNA in Arabidopsis thaliana: Generation of 12,028 non-redundant expressed sequence tags from normalized and size-selected cDNA libraries
DNA Res. 7, 175-180 (2000)
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/strain="Columbia"
/db_xref="taxon:3702"
/clone="Apx20903_f"
/clone_lib="Arabidopsis thaliana Above-ground
two to six-week old plants Columbia"
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/dev_stage="two to six-week old plants"
/note="Vector: pBluescriptII SK-; Site_
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/strain="Columbia"
/db_xref="taxon:3702"
/clone="APZ32g06F"
 /note="Vector:
                 /tissue_type="aboveground organs"
/dev_stage="two to six-week old"
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A1995637
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TGTTGGTTCACAAGATTTGACA 233
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Arabidopsis thaliana Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis. 1 (bases 1 to 530)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Fax: 3
Email:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Genome Systems, Inc., a wholly owned subsidiary of Incyte Pharmaceuticals, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: David Smoller, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Arabidopsis thaliana Gene Expression MicroArray Unpublished (1999)
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Location/Qualifiers
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                                                                                                                                                                        weeks. Plants were grown in 1:1:1 peat moss/vermiculite/perlite soil at 22 deg. C +/- 3 deg. under constant light, and watered with fertilizer. cD synthesis was initiated using a NotI-oligo(dT) primer Double-stranded cDNA was blunted, ligated to SalI adap digested with NotI, size-selected, and cloned into NotI and SalI sites of the psPORT vector."
                                                                                                                                                                                                                                                                                                                  /Clone_lib="A. thaliana, Columbia Col-0, inflorescence-1"
/tissue_type="inflorescence"
/dev_stage="4 - 7 weeks"
/note="vector: psporT; Site_1: NotI; Site_2: SalI; cDNA
library was derived from untreated inflorescence tissue
from Arabidopsis thaliana, Columbia Col-0, at 4 - 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /organism="Arabidopsis thaliana"
/cultivar="Columbia Col-0"
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Lsmatches 2;
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No. 1.1e+02;
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                                                                                                                                                                                          TTGGATCCCAAGATTTGATA 55
CNS02BWA 934 bp DNA GSS 12-MAY-2000 Tetraodon nigroviridis genome survey sequence PUC-Ori end of clone 254C22 of library G from Tetraodon nigroviridis, genomic survey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Laboratory of Cellular and Developmental Biology NIDDK, National Institutes of Health 6 Center Drive MSC 2715, Bldg 6, Rm B1-13, Bether Fax: (301) 496 5239
Email: oliver@helix.nih.gov,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   http://www.niddk.nih.gov/intram/people/boliver.htm Tissue isolation and library construction performed at the National Institute of Diabetes and Digestive and Kidney Diseases, NIH (see http://www.niddk.nih.gov/intram/people/boliver.htm). DNA sequencing and analyses performed by National Institutes of Health Intramural Sequencing Center (NISC; see http://www.nisc.nih.gov).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             bs66h08.y1 Drosophila melanogaster adult testis library Drosophila melanogaster cDNA clone bs66h08 5', mRNA sequence.
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Contact: Brian Oliver
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primer: M13RP1 reverse primer (ABI).
Location/Qualifiers
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                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                            /note="Organ: testis; Vector: pBlueScript SK (Stratagene);
/note="Organ: testis; Vector: pBlueScript SK (Stratagene);
Site_1: EcoR I; Site_2: Xho I; Testes dissected from 1-5
day adult y[*] w[67c1]/Y males raised at 25cC. RNA
isolated using Trizol (Life Technologies) and a single
round of Poly(A)+ selection using Oligotex (Qiagen). cDNA
library constructed using Stratagene ZAP-cDNA syntesis
kit. Oligo dT-primed, size fractionated -1-6 kb, and
directionally cloned at EcoRI and XhoI in Un1-ZAP XR.
Following a single round of amplification pBlueScript SK
phagemids were mass excised. A distribution channel for
clones is being sought, but not currently available.
Requests for clones cannot be honored."
73 a 146 c 161 g 128 t
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/strain="y[*] w[67cl]/Y"
/db_xref="taxon:7227"
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/lab_host="SOLR (Stratagene)"
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                                                                                                                                                                                                                                                                                                                                                                                                           sequence.
AL190387.1 GI:7828491
AL190387.1 GSS; genome survey sequence.
Tetraodon nigroviridis.
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Direct Submission
Submitted (12-APR-2000) to the EMBL/GenBank/DDBJ databases
Submitted (12-APR-2000) to the EMBL/GenBank/DDBJ databases
This sequence is a single read and was generated as part of a large
scale clone-end sequencing project of the Tetraodon nigroviridis
For more information, please take a look at
                                                                                                          Medicago Truncatula
Medicago truncatula
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Epermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
Rosidae; eurosids I; Fabales; Fabaceae; Papilionoideae; Trifolieae
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7 (bases 1 to 934)

8 (Bases 1 to 934)

1 (Bases 1 to 934)
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http://www.genoscope.cns.fr/Tetraodon.
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Roest-Crollius,H., Jaillon,O., Dasilva,C., Bouneau,L., Fisher,C.,
Bernot,A., Fizames,C., Wincker,P., Brottier,P., Quetier,F.,
Saurin,W. and Weissenbach,J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Charaterization and repeat analysis of the compact genome of the freshwater pufferfish Tetraodon nigroviridis
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Unpublished (2000)
Contact: Deborah P
                                                    Fedorova, M., Pierson, B.L., Gonzales, M.B. and Ellis, L.
                                                                                                                                                                                                                                                              sequence.
BF006444
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                                    from Medicago truncatula
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PUC-Ori"
175 c 207 g 259
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /organism="Tetraodon nigroviridis"
/db_xref="taxon:99883"
/clone="254C22"
/clone_lib="G"
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                                                                                                                                                                                                                                                                                                   truncatula
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                                                                       Samac, D.A., Gantt, J.S.,
                                      leaves
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                                      and cotyledons
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                                                                                                                                                                                                                                                                                  AUTHORS
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Ffax: 651 649 5058
Email: debbys@puccini.crl.umn.edu
University of Minnesorta name: M275482e TIGR
MTLBA56TK More information is available at:
http://chrysie.tamu.edu/medicago
Seq primer: SKmod (CTA 9AA CTA 9tg 9AT CC).
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Department of Plant Pathology
University of Minnesota
495 Borlaug Hall, 1991 Upper
Email: debbys@puccini.crl.umn.edu
University of Minnesota name: MZ75292e TIGR
MTLAU47TK More information is available at:
http://chrysie.tamu.edu/medicago
Seg primer: SKmod (CTA gAA CTA gtg gAT CC).
                                                                                                 495 Borlaug Hall, 1991 Upper Buford
Tel: 612 625 1243
Fax: 651 649 5058
                                                                                                                                                                                                                                       Fedorova,M., Pierson,B.L., Samac,D.A., Gantt,J.S., Vance,C.P., Gonzales,M.B. and Ellis,L.
ESTs from Medicago truncatula leaves and cotyledons
                                                                                                                                                                                                                                                                                                                                          Medicago truncatula

Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;

Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;

Rosidae; eurosids I; Fabales; Fabaceae; Papillonoldeae; Trifolieae;
                                                                                                                                                                                                                                                                                                                                                                                                                                         BF006254.1
EST.
                                                                                                                                                           Department of Plant Pathology University of Minnesota
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                                                                                                                                                                                                  Contact: Deborah A. Samac
                                                                                                                                                                                                                      Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                         barrel medic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note-"vector: pBluescript SK +/-; Site_1: EcoRI; Site_2: XhoI; cDNA was prepared from polyA+ enriched RNA from the mixture of cotyledons of five days old plants and leaves of two weeks old plants. The cDNA was directionally ligated into the Uni-ZAP XR vector from Stratagene and packaged using Gigapack III Gold packaging extracts. Plasmids containing cDNA inserts were excised from the recombinant lambda-ZAP phage using Ex-Assist helper phage and propagated in SOLR cells."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /dev_stage="mixture of cotyledons from five plants and leaves obtained from two weeks ol /lab_host="E. coli strain SOLR"
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/cultivar="genotype A17"
/db_xref="taxon:3880"
/clone="pDSLC-41J15"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BB288727 286 bp. mRNA
BB288727 RIKEN full-length enriched,
Clone B020032C12 3', mRNA sequence.
BB288727
                                                                                                                                                                                                                                                                                                                                                                                                                                                       P., Endo, T., Fukuda, S., Fukunishi, Y., Hara, A., Hayatsu, N., Hiro, A., Hayatsu, N., Fukuda, S., Fukunishi, Y., Ishikawa, T., Ishikawa, T., Ishikawa, T., Ishikawa, T., Kalota, K., Kagawa, I., Kal, C., Kawal, J., Kikuchi, N., Kiyosawa, H., Kojima, Y., Kondo, S., Koya, S., Kurihara, C., Kusakabe, M. Matsuyama, T., MIKi, R., Mizuno, Y., Nakamura, M., Odda, H., Okazaki, Y., Ono, T., Owa, C., Saito, H., Sakai, C., Sato, K., Shibata, Y., Shibata, Y., Shipata, Y., Shibata, Y., Shipata, Y., Shipata, Y., Shipata, Y., Shigamara, Y., Shigamara, Y., Shigamara, Y., Shigamara, Y., Shigamara, A., Takahashi, F., Tominaga, N., Toya, T., Tsunoda, Y., Watahiki, A., Watanabe, S., Yamamura, T., Yamanaka, I., Yano, R., Yasunishi, A., Yokota, T., Yoshida, K., Yoshiki, A., Yoshino, M., Muramatsu, M., and Hayashizaki, Y.
                                                                                                                                                                                 Laboratory for Genome Exploration Research Group, RIKEN Genomic Sciences Center(GSC), Yokohama Institute
The Institute of Physical and Chemical Research (RIKEN)
1-7-22 Suehiro-cho, Tsurumi-ku, Yokohama, Kanagawa 230-0045, Japanel: 81-45-503-9222
Fax: 81-45-503-9216
                      Email: genome-res@gsc.riken.go.jp,
URL:http://genome.gsc.riken.go.jp/
Carninci,P., Nishiyama,Y., Westover,A., Itoh,M., Nagaoka,S., Sasaki,N., Okazaki,Y., Muramatsu,M. and Hayashizaki,Y.
Thermostabilization and thermoactivation of thermolabile enzymes by
                                                                                                                                                                                                                                                                                                                                                                         Unpublished (2000)
Contact: Yoshihide Hayashizaki
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 286)
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                                                                                                                                                                                                                                                                                                                                                                                                                                          RIKEN Mouse ESTs (Konno, H., et al.)
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//Clone_lib="DSLC"
//tissue_type="leaves and cotyledons from five days old
//dev_stage="mixture of cotyledons from five days old
//dev_stage="mixture of cotyledons from five days old
plants and leaves obtained from two weeks old plants"
//lab_host="E. coli strain SOLR"
//note="Vector: pBluescript SK +/-; Site_1: EcoRI; Site_2:
//note="vector: pBluescript Sold RNA from the
mixture of cotyledons of five days old plants and leaves
of two weeks old plants. The cDNA was directionally
ligated into the Uni-ZAP XR vector from Stratagene and
packaged using Gigapack III Gold packaging extracts.
Plasmids containing cDNA inserts were excised from the
recombinant lambda-ZAP phage using Ex-Assist helper phage
and propagated in SOLR cells."
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/cultivar="genotype A17"
/db_xref="taxon:3880"
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. No. 3.2e+02;
ismatches 2;
                                                                                                                                                                                                                                                  Yokohama, Kanagawa 230-0045, Japan
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synthesis of full length
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RESULT 11
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                                                                                                                 McCarter,J., Clifton,S., Chiapelli,B., Pape,D., Martin,J., Wylie,T., Dante,M., Marra,M., Hillier,L., Kucaba,T., Theising,B., Bowers,Y., Gibbons,M., Ritter,E., Bentett,J., Franklin,C., Tsagareishvili,R., Ronko,I., Kennedy,S., Maguire,L., Beck,C., Underwood,K., Steptoe,M., Allen,M., Person,B., Swaller,T., Harvey,N., Schurk,R., Kohn,S., Shin,T., Jackson,Y., Cardenas,M., McCann,R., Waterston,R. and
The Washington Univ. Nematode EST Project, 1999 Washington University School of Medicine
                                     The Washington Univ. Nematode EST Project, 1999
Unpublished (1999)
Contact: McCarter JP
                                                                                                                                                                                                                                                                 Eukaryota: Metazoa; Nematoda; Chromadorea; Rhabditida;
Panagrolaimoidea; Strongyloididae; Strongyloides.
                                                                                                                                                                                                                                                                                                           Strongyloides stercoralis. Strongyloides stercoralis
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Automated filtration-based high-throughput plasmid preparation
Automated filtration-based 1999)
system. Genome Res. 9 (5), 463-470 (1999)
Carninci,P. and Hayashizaki,Y.
High-efficiency full-length cDNA cloning. Methods Enzymol. 303,
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                                                                                                                                                                                                                                           (bases, 1 to 407)
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/db_xref="taxon:10090"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tel: 314 286 1800
Fax: 314 286 1810
Email: est@watson.wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18;
Email: jwallace@u.washington.edu

Email: jwallace@u.washington.edu

Clones are derived from the human BAC library RPCI-11. For BAC

Clones are derived from the human BAC library availability, please contact Pieter de Jong

library availability, please contact Pieter de Jong

(pleter@dejong.med.buffalo.edu). Clones may be purchased from

(pleter@dejong.med.buffalo.edu). Clones may be purchased from

BACPAC Resources (http://bcac.ned.buffalo.edu/ordering_bac.htm)

or from Resear h Genetics (info@resgen.com). BAC end Web Server:

http://www.htsc.washington.edu

plate: 642 row: J column: 9
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1 (bases 1 to 529)
Mahairas, G.G., Wallace, J.C., Smith, K., Swartzell, S., Holzman, T., Mahairas, G.G., Wallace, J.C., Smith, K., Swartzell, S., Adams, M.D. Keller, A., Shaker, R., Furlong, J., Young, J., Zhao, S., Adams, M.D.
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                                                                                                                                                                                                            Contact: Mahairas GG, Wallace JC, Hood L
High Throughput Sequencing Center
University of Washington
401 Queen Anne Avenue North, Seattle, WA 98109, USA
Tel: (206) 616-3618
Fax: (206) 616-3887
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(range, 100-1700)."

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(Stratagene); Site_1: // (Stratagene); Site_1: // (Note="Wector: Lambda Uni-ZAP XR (Stratagene); X 10E3 ECORI; Site_2: XhOI; mRNA was purified from 2 x 10E3 Frabditionally infected was purified from gerbils infected with larvae originally isolated experimentally infected dogs. cDNA was constructed and, using adaptors, was cloned unidirectionally into the and unample from the EcoRI site to the XhOI site. The library has an unamplified titer of 1 x 10E5 pfu/ml and an amplified, undiluted titer of 9 x 10E11 pfu/ml. The average insert size of the unamplified library is 675 bp (range, 100-1700)."

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/strain="Rhabditiform larvae obtained from gerbils"
/db_xref."taxon:6248"
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Class: BAC ends
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                                                                                                                                                                                                                                                                                                                                                                                                                     Expressed Sequence Tags from the Samuel Roberts Noble Foundation Medicago truncatula stem library Unpublished (2000) on Apr 14, 2000 this sequence version replaced gi:7569784. Contact: Dixon RA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 barrel medic. Medicago truncatula Medicago truncatula Embryophyta; Embryophyta; Tracheophyta; Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Spermatophyta; Magnoliophyta; Fabales; Fabaceae; Papilionoideae; Trifolieae
                                                                                                                                                                                                                                                                                                                The Samuel Roberts Noble Foundation 2510 Sam Noble Parkway, Ardmore, OK 73402, USA Tel: 580 221 7302 Fax: 580 221 7380
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 (bases 1 to 646)
He,x.-z., Shadle,G., Scott,A.D., Harris,A.R.,
,C.J., Flores,H.R., Inman,J.T., Weller,J.W.,
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                                                                                                                                                                                                                Email: radixon@noble.org
Insert Length: 674 Std Error: 0.00
Plate: 082 row: E column: 04
Seq primer: TCACACAGGAAACAGCTATGAC.
Location/Qualifiers
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/organism="Homo sapiens"
/db xref="taxon:9606"
/clone="Plate=642 Col=9 Row=J"
/clone_lib="RPCI-11 Human Male BAC Library"
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    /clone_lib="Developing stem"
/tissue_type="stem"
/dev_stage="pooled developmental"
/note="Vector: Lambda Zap; Contains a mixture
internodal stem segments"
142 c 129 g 195 t
                                                                                                                               /organism="Medicago truncatula"
/db_xref="taxon:3880"
/clone="NF082E04ST"
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мау,G.D. and Dixon
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DU448853 671 bp mRNA EST 16-MAR-2001
NF003C10INIF1082 Insect herbivory Medicago truncatula cDNA clone
BC448853 BC448853 BC448853.1 GI:13367634
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               barrel medic.

Medicago truncatula

Medicago; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;

Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;

Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;

Rosidae; eurosids I; Fabales; Fabaceae; Papilionoideae; Trifolieae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 (bases 1 to 671) ... Harris, A.R., Gonzales, R.A., Bell, C.J., Flores, H.R., Inman, J.T., Weller, J.W. and May, G.D. Expressed Sequence Tags from the Samuel Roberts Noble Foundation Medicago truncatula insect herbivory library Unpublished (2000) Contact: Korth K Dept. of Plant Pathology University of Arkansas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Email: kkorth@comp.uark.edu
Insert Length: 671 Std Err
Plate: 003 row: C column:
Seg primer: TCACACAGGAAACAGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               217 Plant Science Building, Fayetteville, AR 72701, USA Tel: 501 575 5191
Fax: 501 575 7601
                                mRNA sequence.
                                               BE283043 762 bp mRNA
601101323F1 NCI_CGAP_Lu29 Mus
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                                                                                                                                                                                                                                                                                                                                      /Clone_lib="Insect herbivory"
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/dev_stage="mature"
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/note="Vector: Lambda Zap; Library was produced from fully
expanded M. truncatula leaves of plants fed upon by
spodoptera exigua (beet armyworm) for 24 hours. Systemic
(undamaged leaves from injured plants) and wounded leaves
were harvested and pooled."
a 151 c 131 g 203 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /clone="NF003C10IN"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /organism="Medicago truncatula"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        .671
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94.78;
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94.78;
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                                                                                                                                                                                                                                     0;
                                                                                                                                                                                                                                     Score 17.4; DB 11;
Pred. No. 5e+02;
0; Mismatches 1;
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                                                     musculus
                                                     cDNA clone
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IMAGE:3493829
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Search completed: December 15, Job time: 4977 sec

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REFERENCE
AUTHORS
TITLE
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                                                                                                        Query Match
Best Local 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    source
                    1 tgttggatcacaagatttg
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Email: cgapbs r@mail.nih.gov
Tissue Procurement: Gilbert Smith, Ph.D.

CDNA Library Preparation: Life Technologies, Inc.

cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)

DNA Sequencing by: Incyte Genomics, Inc.

Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mus musculus Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Eukaryota; Metazoa; Chordata; Sciurognathi; Muridae; Musinae; Mus. Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Musinae; Mus. 1 (bases 1 to 762)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC) Unpublished (1999)
                                                                                    l Similarity
18; Conserv
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plate: LLAM8541 row: n column: 06
High quality sequence stop: 661.
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house mouse
                                                                                                                                                                                                                     193
                                                                                    Conservative
                                                                                                                                                                                                               /note="Organ: lung; Vector: pCMV-SPORT6; Site_1: SalI; Site_2: NotI; Cloned unidirectionally. Primer: Oligo dT. Library constructed by Life Technologies. Investigator providing samples: Gilbert Smith, NIH"

a 172 c 207 g 190 t
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94.7%;
                                        19
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                                                                                                          Score 17.4; DB 10
Pred. No. 5.1e+02;
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                                                                                                                             DB 10;
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Page 8

Run

on:

OM nucleic - nucleic search, using sw model

GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd

December 15, 2001, 05:42:56;

(without alignments)
15.407 Million cell updates/sec Search time 7962.77 Seconds

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Database :
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 2000000000
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Perfect score:
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          US-09-380-826A-5
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Gapop 10.0 , Gapext 1.0
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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Sequence 60, Appli Sequence 8, Appli Sequence 9, Appli Sequence 10, Appl Sequence 655, App Sequence 656, App Sequence 90, Appl	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0	7, API 7, API 4508, 4785, 186, 40, AI	e 7517, e 7816, e 10702, e 10703, e 11663, e 11664, e 11665, e 12098, e 12170, e 12170, e 4734, P	Sequence 5, Appli Sequence 1005, Ap Sequence 1005, Ap Sequence 1287, Ap Sequence 1793, Ap Sequence 6568, Ap Sequence 7108, Ap Sequence 7108, Ap

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100 100 100

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> US-08-584-040-8444 US-08-585-684-113 US-08-585-684-114

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US-09-708-690-4100

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APPLICATION NUMBER: 1905494/97
PRIOR APPLICATION NUMBER: 05494/97
PRIOR APPLICATION NUMBER: 05494/97
PRIOR APPLICATION NUMBER: 05494/97
PRIOR APPLICATION NUMBER: 075494/97
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PRIOR FI
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US-09-380-826A-5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILE REFERENCE: 68126881206940
CURRENT APPLICATION NUMBER: PCT/US99/13800
CURRENT FILING DATE: 1999-06-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Genzyme Corporation
APPLICANT: Roberts, Bruce L.
APPLICANT: Shankara, Srinivas
TITLE OF INVENTION: PREPARATION AND USE OF SUPERIOR VACCINES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EARLIER APPLICATION NUMBER: 60/090,039
EARLIER FILING DATE: 1998-06-19
EARLIER APPLICATION NUMBER: 60/090,040
EARLIER FILING DATE: 1998-06-19
EARLIER APPLICATION NUMBER: 60/090,041
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ORGANISM: Artificial Sequence
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ER APPLICATION NUMBER: 60/090,035
ER FILING DATE: 1998-06-19
ER APPLICATION NUMBER: 60/089,993
ER FILING DATE: 1998-06-19
ER APPLICATION NUMBER: 60/089,992
ER APPLICATION NUMBER: 60/089,992
ER APPLICATION NUMBER: 60/089,092
ER APPLICATION NUMBER: 60/090,072
ER FILING DATE: 1998-06-19
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APPLICATION NUMBER: 60/090,039
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2.5e+09;
hes 0;
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Sequence 1287, Application US/09335032

GENERAL INFORMATION:
APPLICANT: Velculescu, Victor
APPLICANT: Vogelstein, Bert
APPLICANT: Kinzler, Kenneth
TITLE OF INVENTION: Characterization of the Yea
TITLE OF INVENTION: Transcriptome
FILE REFERENCE: 01107.78572
CURRENT APPLICATION UNMBER: US/09/335,032
CURRENT EILING DATE: 1999-06-16
PRIOR APPLICATION NUMBER: US 09/015,917
PRIOR FILING DATE: 1997-01-23
PRIOR APPLICATION NUMBER: US 09/012,031
PRIOR APPLICATION NUMBER: US 09/012,031
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                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
Theshes 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; TYPE: DNA; ORGANISM: Homo sapiens PCT-US99-13800-1005
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                                                                                                                                                                                                                                                                            US-09-335-032-1287/c
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SOFTWARE: FastSEQ for Windows Version
SEQ ID NO 1005
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EARLIER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     EARLIER FILING DATE: 1998-12-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     EARLIER FILING DATE: 1998-06-19
EARLIER APPLICATION NUMBER: 60/111,715
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ER FILING DATE: 1998-06-19
ER APPLICATION NUMBER: 60/090,078
ER FILING DATE: 1998-06-19
ER APPLICATION NUMBER: 60/090,047
ER FILING DATE: 1998-06-19
ER FILING DATE: 1998-06-19
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R FILING DATE: 1998-06-19
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FILING DATE: 1998-06-19
APPLICATION NUMBER: 60/089,994
FILING DATE: 1998-06-19
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FILING DATE: 1998-06-19
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Pred. No. 3.3
); Mismatches
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3.3e+05;
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; TYPE: DNA; ORGANISM: Saccharomyces cerevisiae US-09-335-032-1287

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SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 1287 LENGTH: 10

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Query Match
Best Local Similarity
7; Conserve
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US-09-335-032-1793/c
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Best Local Similarity 100.
Thes 7; Conservative
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1793
                                                     SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 4831
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
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                                                                                        PRIOR APPLICATION NUMBER: US 60/035,917
PRIOR FILING DATE: 1997-01-23
PRIOR APPLICATION NUMBER: US 09/012,031
PRIOR FILING DATE: 1998-01-22
NUMBER OF SEQ ID NOS: 12219
                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/335,032
CURRENT FILING DATE: 1999-06-16
                                                                                                                                                                                                                    APPLICANT: Velculescu, Victor
APPLICANT: Vogelstein, Bert
APPLICANT: Kinzler, Kenneth
TITLE OF INVENTION: Characterization of the Yeast
TITLE OF INVENTION: Transcriptome
FILE REFERENCE: 01107.78572
                                                                                                                                                                                                                                                                                                                                                 sequence 4831, Application US/09335032
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: US 60/035,917
PRIOR FILING DATE: 1997-01-23
PRIOR APPLICATION NUMBER: US 09/012,031
PRIOR FILING DATE: 1998-01-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Velculescu, Victor
APPLICANT: Vogelstein, Bert
APPLICANT: Kinzler, Kenneth
TITLE OF INVENTION: Characterization of the Yeast
TITLE OF INVENTION: Transcriptome
FILE REFERENCE: 01107.78572
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/335,032 CURRENT FILING DATE: 1999-06-16
                                                                                                                                                                                                                                                                                                                              ENERAL INFORMATION:
ORGANISM: Saccharomyces cerevisiae
                                        ENGTH:
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Pred. No.
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3.3e+05;
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Query Match Best Local Similarity

100.0%;

Score 7; Pred. No.

DB 17; 3.3e+05;

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PRIOR APPLICATION NUMBER: US 09/012,031
PRIOR FILING DATE: 1998-01-22
NUMBER OF SEQ ID NOS: 12219
SOPTWARE: FRASTSEQ for Windows Version 4.0
SEQ ID NO 6568
LENGTH: 10
TYPE: DNA
ORGANISM: Saccharomyces cerevisiae
US-09-335-032-6568
                                                                                                                                                                                                                                                                       US-09-335-032-7108/c
Sequence 7108, Application US/09335032
GENERAL INFORMATION:
APPLICANT: Veiculescu, Victor
APPLICANT: Vogelstein, Bert
APPLICANT: Kinzler, Kenneth
TITLE OF INVENTION: Characterization of the
TITLE OF INVENTION: Transcriptome
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US-09-335-032-7108
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Best Local Similarity
Matches 7; Conser
              NUMBER OF SEQ ID NOS: 12219

SOFTWARE: FRATSEQ for Windows Version 4.0

SEQ ID NO 7108

LENGTH: 10

TYPE: DNA

ORGANISM: Saccharomyces cerevisiae
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Best Local
                                                                                                                                 FILE REFERENCE: 01107.78572
CURRENT APPLICATION NUMBER: US/09/335,032
CURRENT FILING DATE: 1999-06-16
PRIOR APPLICATION NUMBER: US 60/035,917
PRIOR TILING DATE: 1997-01-23
PRIOR APPLICATION NUMBER: US 09/012,031
PRIOR APPLICATION NUMBER: US 09/012,031
PRIOR APPLICATION NUMBER: US 09/012,031
PRIOR FILING DATE: 1998-01-22
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CURRENT FILING DATE: 1999-06-16
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APPLICANT: Vogelstein, Bert
APPLICANT: Kinzler, Kenneth
TITLE OF INVENTION: Characterization of the Yeast
TITLE OF INVENTION: Transcriptome
FILE REFERENCE: 01107.78572
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Pred. No.
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3.3e+05;
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TITLE OF INVENTION: Characterization of the Yeast
TITLE OF INVENTION: Transcriptome.
FILE REFERENCE: 01107.78572
CURRENT APPLICATION NUMBER: US/09/335.032
CURRENT FILING DATE: 1999-06-16
PRIOR APPLICATION NUMBER: US 09/015,917
PRIOR FILING DATE: 1997-01-23
PRIOR APPLICATION NUMBER: US 09/012,031
PRIOR APPLICATION NUMBER: US 09/012,031
PRIOR FILING DATE: 1998-01-22
NUMBER OF SEQ ID NOS: 12219
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 7128
LENGTH: 10
TYPE: DNA
                                                                                                                                                                                                                                 APPLICANT: Vogelstein, Bert
APPLICANT: Vogelstein, Bert
APPLICANT: Winzler, Kenneth
APPLICANT: Kinzler, Kenneth
APPLICANT: Kinzler, Kenneth
TITLE OF INVENTION: Transcriptome
FILE OF INVENTION: Transcriptome
FILE REFERENCE: 01107.78572
CURRENT APPLICATION NUMBER: US/09/335,032
CURRENT APPLICATION NUMBER: US 60/035,917
PRIOR APPLICATION NUMBER: US 60/035,917
PRIOR FILING DATE: 1997-01-23
PRIOR APPLICATION NUMBER: US 09/012,031
PRIOR FILING DATE: 1998-01-22
NUMBER OF SEQ ID NOS: 12219
NUMBER OF SEQ ID NOS: 12219
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 7153
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US-09-335-032-7128
; Sequence 7128, Application US/09335032
; Sequence 7128, Application US/09335032
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US-09-335-032-7153
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                                                                                                                                                                      ; ORGANISM: Saccharomyces cerevisiae US-09-335-032-7153
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                                                       Query Match
Best Local Similarity lov.
7; Conservative
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                                                                             100.0%; Sc
100.0%; Pr
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Pred. No.
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                                                                                   3.3e+05;
hes 0;
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APPLICANT: Vogelstein, Bert
APPLICANT: Kinzler, Kenneth
TITLE OF INVENTION: Characterization of the Yeast
TITLE OF INVENTION: Transcriptome
FILE REFERENCE: 01107.78572
CURRENT APPLICATION NUMBER: US/09/335,032
CURRENT FILING DATE: 1999-06-16
PRIOR APPLICATION NUMBER: US 60/035,917
PRIOR FILING DATE: 1997-01-23
PRIOR APPLICATION NUMBER: US 09/012,031
PRIOR FILING DATE: 1998-01-22
NUMBER OF SEQ ID NOS: 12219
SOFTMARE: FastSEQ for Windows Version 4.0
SEQ ID NO 7517
LENGTH: 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; ORGANISM: Saccharomyces cerevisiae US-09-335-032-7517
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                                                                                                                                                                       ; TYPE: DNA ; ORGANISM: Saccharomyces cerevisiae US-09-335-032-7816
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US-09-335-032-7816
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GENERAL INFORMATION:
APPLICANT: Velculescu, Victor
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Best Local Similarity 100.
Conservative
                                                                                                                                                                                                                              NUMBER OF SEQ ID NOS: 12219
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 7816
LENGTH: 10
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                                                                                          Query Match
Best Local
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APPLICANT: Vogelstein, Bert
APPLICANT: Kinzler, Kenneth
TITLE OF INVENTION: Characterization of the Yeast
TITLE OF INVENTION: Transcriptome
FILE REFERENCE: 01107.78572
CURRENT APPLICATION NUMBER: US/09/335,032
CURRENT FILING DATE: 1999-06-16
                                                                            Matches
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PRIOR FILING DATE: 1997-01-23
PRIOR APPLICATION NUMBER: US 09/012,031
PRIOR FILING DATE: 1998-01-22
                                                                        Local Similarity nes 7; Conserv
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                                    1 tttgata 7
                                                                    100.0%; Score 7; DB 17; ilarity 100.0%; Pred. No. 3.3e+05; Conservative 0; Mismatches 0.
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US-09-335-032-8062/c ; Sequence 8062, Application US/09335032

RESULT

GENERAL INFORMATION:

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RESULT 14
US-09-315-032-11663/c
Sequence 11663, Application US/09335032
GENERAL INFORMATION:
APPLICANT: Velculescu, Victor
APPLICANT: Vogelstein, Bert
APPLICANT: Kinzler, Kenneth
TITLE OF INVENTION: Characterization of the Yeast
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Best Local Similarity
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APPLICANT: Vogelstein, Bert
APPLICANT: Kinzler, Kenneth
TITLE OF INVENTION: Characterization of the Yeast
TITLE OF INVENTION: Transcriptome
FILE REFERENCE: 01107,78572
CURRENT APPLICATION NUMBER: US/09/335,032
CURRENT EILING DATE: 1999-06-16
PRIOR APPLICATION NUMBER: US 60/035,917
PRIOR FILING DATE: 1997-01-23
PRIOR FILING DATE: 1997-01-23
PRIOR FILING DATE: 1998-01-22
PRIOR FILING DATE: 1998-01-23
PRIOR FILING DATE: 1998-01-23
PRIOR FILING DATE: 1998-01-23
PRIOR FILING DATE: 1998-01-29
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Velculescu, Victor
APPLICANT: Vogelstein, Bert
APPLICANT: Kinzler, Kenneth
TITLE OF INVENTION: Characterization of the Yeast
TITLE OF INVENTION: Transcriptome
FILE REFERENCE: 01107.78572
CURRENT APPLICATION NUMBER: US/09/335,032
CURRENT FILING DATE: 1999-06-16
PRIOR APPLICATION NUMBER: US 60/035,917
PRIOR APPLICATION NUMBER: US 09/012,031
PRIOR FILING DATE: 1997-01-22
PRIOR FILING DATE: 1998-01-22
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Best Local Similarity
"~*~hes 7; Conserv?
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US-09-335-032-10702/c
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 10702
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LENGTH: 10
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Pred. No.
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Pred. No.
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hes 0;
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Search completed: December 15, 2001, 05:42:57 Job time: 15822 sec

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APPLICANT: KINZLEY, Kenneth

TITLE OF INVENTION: Characterization of the Yeast

TITLE OF INVENTION: Transcriptome

FILE REFERENCE: 01107.78572

CURRENT APPLICATION NUMBER: US/09/335,032

CURRENT APPLICATION NUMBER: US 60/035,917

PRIOR APPLICATION NUMBER: US 60/035,917

PRIOR FILING DATE: 1999-01-23

PRIOR FILING DATE: 1999-01-23

PRIOR APPLICATION NUMBER: US 09/012,031

PRIOR FILING DATE: 1998-01-22

NUMBER OF SEQ ID NOS: 12219

SOFTMARE: FastSEQ for Windows Version 4.0

SEQ ID NO 11664

LENGTH: 10

TYPE: DNA
ORGANISM: Saccharomyces cerevisiae

US-09-335-032-11664
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/335,032
CURRENT FILING DATE: 1999-06-16
PRIOR APPLICATION NUMBER: US 60/035,917
PRIOR FILING DATE: 1997-01-23
PRIOR APPLICATION NUMBER: US 09/012,031
PRIOR FILING DATE: 1998-01-22
NUMBER OF SEQ ID NOS: 12219
SOFTWARR: FRStSEQ for Windows Version 4.0
SEQ ID NO 11663
LENGTH: 10
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                                                   Query Match
Best Local Similarity
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US-09-335-032-11664/c
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Best Local Similarity
Matches 7; Conserv
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Pred. No.
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. 3.3e+05;
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3.3e+05;
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq
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/cgn2_6/ptodata/1/pna/US07_NEW_COMB.seq:*
/cgn2_6/ptodata/1/pna/US08_NEW_COMB.seq:*
/cgn2_6/ptodata/1/pna/US09_NEW_COMB.seq:*
/cgn2_6/ptodata/1/pna/US09_NEW_COMB.seq:*
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BB
                                       US-09-685-664B-4100
US-09-870-161-4100
US-09-870-161-5694
US-09-8870-161-5694
US-09-685-664B-117
US-09-685-664B-117
US-09-685-664B-883
US-09-685-664B-2418
US-09-685-664B-2418
US-09-685-664B-3110
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21.200 Million cell updates/sec
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4100, Ap
5694, Ap
5694, Ap
117, App
118, App
883, App
883, App
2417, Ap
3108, Ap
3110, Ap
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27 7 100.0 17 6 US-09-670-607-971 28 7 100.0 17 6 US-09-670-607-972 29 7 100.0 17 6 US-09-670-607-972 29 7 100.0 17 6 US-09-670-607-1467 30 7 100.0 17 6 US-09-870-161-116 31 7 100.0 17 6 US-09-870-161-117 32 7 100.0 17 6 US-09-870-161-118 34 7 100.0 17 6 US-09-870-161-188 35 7 100.0 17 6 US-09-870-161-883 36 7 100.0 17 6 US-09-870-161-883 37 100.0 17 6 US-09-870-161-2418 38 7 100.0 17 6 US-09-870-161-2108 39 7 100.0 17 6 US-09-870-161-31108 39 7 100.0 17 6 US-09-870-161-31108 30 7 100.0 17 6 US-09-870-161-31108 31 7 100.0 17 6 US-09-870-161-3119 40 7 100.0 17 6 US-09-870-161-3119 41 7 100.0 17 6 US-09-870-161-3119 42 7 100.0 17 6 US-09-870-161-3119 43 7 100.0 17 6 US-09-870-161-3119 44 7 100.0 17 6 US-09-870-161-3119 45 7 100.0 17 6 US-09-870-161-3129 46 7 100.0 17 6 US-09-870-161-3129	G	G		0						C									C	
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US-09-670-607-971 US-09-670-607-972 US-09-670-607-1468 US-09-670-607-1468 US-09-870-161-116 US-09-870-161-117 US-09-870-161-1882 US-09-870-161-883 US-09-870-161-2418 US-09-870-161-2418 US-09-870-161-3108 US-09-870-161-3110 US-09-870-161-3110 US-09-870-161-3110 US-09-870-161-3196 US-09-870-161-3196 US-09-870-161-3198	6	σ	6	σ	σ	σ	σ	6	σ	σ	σ	σ	σ	σ	σ	σ	σ	σ	6	(
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971, App 972, App 972, App 1168, App 1116, App 1118, App 118, App 2417, App 2417, App 31109, Ap 31109, Ap 31100, Ap 31101, Ap	Ą	Ą	Αp	Αp	Ąρ	Ąρ	Ąρ	Ąρ	Αp	Ąρ	qq	ďď	đď	ďď	ďď	Αp	Ąρ	ď	qq	Ċ

ALIGNMENTS

US-09-685-664B-4100/c Sequence 4100, Application US/09685664B GENERAL INFORMATION: APPLICANT: Ribozyme Pharmaceuticals, Inc. APPLICANT: Pavco, Pam APPLICANT: McSwiggen, Jim APPLICANT: Stinchcomb, Dan APPLICANT: Escobedo, Jaime В ; TYPE: RNA; Homo sapiens US-09-685-664B-4100 Query Match Best Local S Matches 7 SEQ ID NO 4100 LENGTH: 15 APPLICANT: Stinchcomb, Dan APPLICANT: Escobedo, Jalme TITLE OF INVENTION: Method and Reagent for Treatment of Diseases or Conditions Rel TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Recptor TILE REFERENCE: MBHB00-876-K (400/021) CURRENT APPLICATION NUMBER: US 00/085,664B CURRENT FILING DATE: 2000-10-10 PRIOR APPLICATION NUMBER: US 60/005,974 PRIOR FILING DATE: 1995-10-26 PRIOR APPLICATION NUMBER: US 08/584,040 PRIOR FILING DATE: 1996-01-08 ü 1 tttgata 7 TTTGATA 7 Similarity 7; Conserv Conservative 100.0%; 0, Score 7; DB Pred. No. 7.1 0; Mismatches DB 6; Le 7.1e+04; hes 0; Length 15; 0 Gaps 0,

Sequence 4100, Application US/09870161
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Pavco, Pam
APPLICANT: McSwiggen, Jim

APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions

US-09-870-161-4100/c

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Query Match
Best Local Similarity
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ORGANISM: Homo sapiens
US-09-870-161-5694
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US-09-870-161-5694/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/
CURRENT FILING DATE: 2001-08-2
NUMBER OF SEQ ID NOS: 20821
SOFTWARE: PATENTIN VETSION 3.0
SEQ ID NO 4100
LENGTH: 15
                                                SEQ ID NO 5694
LENGTH: 16
TYPE: RNA
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GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Usman, Nassim
APPLICANT: WCSwiggen, Jim
APPLICANT: MCSwiggen, Jim
TITLE OF INVENTION: Method and Reagent for the Inhibition of Protein-Tyrosine Phospha
TITLE OF INVENTION: (Ptp.1b) Enzyme
                                                                                                                                                                                                                                                                                                                                                                 Sequence 5694, Application US/09870161 GENERAL INFORMATION:
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Best Local Similarity
Matches 3; Conserv
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                                                                                                         TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor FILE REFERENCE: MBHB00-876-M (400/026) CURRENT APPLICATION NUMBER: US/09/870,161 CURRENT FILING DATE: 2001-08-27 NUMBER OF SEQ ID NOS: 20821 SOFTWARE: Patentin version 3.0
                                                                                                                                                                                                                                                                                                      APPLICANT: Ribozyme Pharmaceuticals, Inc APPLICANT: Pavco, Pam APPLICANT: McSwiggen, Jim''.
                                                                                                                                                                                                                                                              APPLICANT: Stinchcomb, Dan APPLICANT: Escobedo, Jaime
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13 TTTGATA 7
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McSwiggen, Jim ': .
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42.9%; Pred. No. 7
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Pred. No. 7.1e+04;
0; Mismatches 0;
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APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method and Reagent for Treatment of Diseases or Conditions Rel
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Recptor
FILE REFERENCE: MBHB00-876-K (400/021)
CURRENT APPLICATION NUMBER: US/09/685,664B
CURRENT APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-10-26
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR APPLICATION NUMBER: US 09/371,772
PRIOR FILING DATE: 1996-01-08
PRIOR APPLICATION NUMBER: US 09/371,772
PRIOR FILING DATE: 1999-08-01
SEQ ID NO 116
SEQ ID NO 116
LENGTH: 17
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; Sequence 116, Application US/09685664B

; GENERAL INFORMATION:
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Best Local Similarity
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Best Local S
Matches 7
; ORGANISM: Homo sapiens US-09-685-664B-116
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APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: M8H800-876-M (400/026)
CURRENT APPLICATION NUMBER: US/09/870,161
CURRENT FILING DATE: 2001-08-27
NUMBER OF SEQ ID NOS: 20821
SOFTWARE: Patentin version 3.0
SEQ ID NO 5978
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TYPE: RNA
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Similarity 100.0%;
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Pred. No.
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7.1e+04;
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Score

DB 6;

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SEQ ID NO 117
LENGTH: 17
TYPE: RNA
ORGANISM: Homo sapiens
US-09-685-664B-117
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Best Local Similarity
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                                                                             PRIOR FILING DATE: 1999-08-10, NUMBER OF SEQ ID NOS: 8231 SOFTWARE: PatentIn version 3.0 SEQ ID NO 118
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ORGANISM: Homo sapiens -09-685-664B-118
                                                                                                                                                                      APPLICANT: Stinchomb, Dan
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method*and Reagent for Treatment of Diseases or Conditions Relate
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Recptor
FILE REFERENCE: MBHB00-876-k (400/021)
CURRENT APPLICATION NUMBER: US/09/685,664B
CURRENT FILING DATE: 2000-10-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-10-26
PRIOR FILING DATE: 1995-10-08
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR FILING DATE: 1996-01-08
PRIOR APPLICATION NUMBER: US 09/371,772
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PRIOR FILING DATE: 1995-10-26
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR FILING DATE: 1996-01-08
PRIOR APPLICATION NUMBER: US 09/371,772
PRIOR FILING DATE: 1999-08-10
NUMBER OF SEQ ID NOS: 8231
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APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method and Reagent for Treatment of Diseases or Conditions Relate
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Recptor
TILE REFERENCE: MBHB00-876-K (400/021)
CURRENT EPPLICATION NUMBER: US/09/685,664B
CURRENT FILING DATE: 2000-10-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Ribozyme Pharmaceuticals, Inc.
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                                                               LENGTH: 17
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Stinchcomb, Dan
Escobedo, Jaime
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); Mismatches 0;
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DB 1;
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TITLE OF INVENTION: Method and Reagent for Treatment of Diseases or Conditions Rel TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Recptor FILE REFERENCE: MBHB00-876-K (400/021)
CURRENT APPLICATION NUMBER: US/09/685,664B
CURRENT FILING DATE: 2000-10-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 60/005,774
PRIOR APPLICATION NUMBER: US 09/371,772
PRIOR APPLICATION NUMBER: US 09/371,772
PRIOR FILING DATE: 1996-01-08
PRIOR APPLICATION NUMBER: US 09/371,772
PRIOR FILING DATE: 1999-08-10
NUMBER OF SEQ ID NOS: 8231
SOFTWARE: PatentIn version 3.0
SEQ ID NO 882
US-09-685-664B-882
APPLICANT: ESCODEDO, Jaime: APPLICANT: ESCODEDO, Jaime: TITLE OF INVENTION: Method and Reagent for Treatment of Diseas: TITLE OF INVENTION: Levels of Vascular Endothelial Growth Fac FILE REFERENCE: MBHB00-876-K (400/021)
CURRENT FILING DATE: 2000-10-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-10-26
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR FILING DATE: 1995-10-08
PRIOR APPLICATION NUMBER: US 09/371,772
PRIOR FILING DATE: 1996-01-08
PRIOR FILING DATE: 1999-08-10
NUMBER OF SEQ ID NOS: 8231
SOFTWARE: Patentin version 3.0
SEQ ID NO 883
LENGTH: 17
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; Sequence 882, Application US/09685664B
; GENERAL INFORMATION:
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Matches 7; Conserv
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APPLICANT: Pavco, Pam
APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Escobedo, Jaime
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; TYPE: RNA
; ORGANISM: Homo sapiens
US-09-685-664B-883
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US-09-685-664B-2417/c
; Sequence 2417, Application US/09685664B
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; TYPE: RNA
; ORGANISM: Mus musculus
US-09-685-664B-2417
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APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Pavco, Pam
APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
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APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Pavco, Pam
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                    APPLICANT: Pavco, Pam
APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method and Reagent for Treatment of Diseases or Conditions Relate
TITLE OF INVENTION: Method and Reagent for Treatment of Escotor Recptor
FILLE REFERENCE: MBHB00-876-K (400/021)
CURRENT FILING DATE: 1000-10-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-01-08
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Stinchcomb, Dan
Escobedo, Jaime
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APPLICANY: Stinchcomb, Dan
APPLICANY: Escobedo, Jaime
APPLICANY: Escobedo, Jaime
TITLE OF INVENTION: Method and Reagent for Treatment of Diseases or Conditions Rel
TITLE OF INVENTION: Method and Reagent for Treatment of Diseases or Conditions Rel
TITLE OF INVENTION: Method and Reagent for Treatment of Diseases or Conditions Rel
TITLE OF INVENTION: Method and Reagent for Treatment of Diseases or Conditions Rel
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TITLE OF INVENTION: Method and Reagent for Treatment of Treatment of Conditions Rel
TITLE OF INVENTION: Method and Reagent for Treatment of Conditions Rel
TREATMENT APPLICATION NUMBER: US 00/005,974
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR APPLICATION NUMBER: US 09/371,772
PRIOR APPLICATION NUMBER: US 09/371,772
PRIOR TILING DATE: 1995-01-08
PRIOR FILLING DATE: 1999-08-10
NUMBER OF SEQ ID NOS: 8.231
SOFTWARE: Patentin version 3.0
SEQ ID NO 3108
LENGTH: 17
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; ORGANISM: Mus musculus
US-09-685-664B-2418
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US-09-685-664B-3109
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Best Local Similarity
Matches 7; Conserv
                                                                                                                                                                                                        Sequence 3109, Application US/09685664B
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Pavco, Pam
APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Escobedo, Jaime
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TITLE OF INVENTION: Method and Reagent for Treatment of Diseases or Conditions TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Recptor FILE REFERENCE: MBHB00-876-K (400/021)
CURRENT APPLICATION NUMBER: US/09/685.664B
CUGRENT FILING DATE: 2000-10-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR PILING DATE: 1995-10-26
PRIOR FILING DATE: 1995-01-08
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7.1e+04;
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Sequence 3110, Application US/0968564B
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Roswiggen, Jim
APPLICANT: McSwiggen, Jim
APPLICANT: Escobedo, Jaime
ITILE OF INVENTION: Method and Reagent for Treatment of Diseases or Conditions Relate
TITLE OF INVENTION: Method and Reagent for Treatment of Diseases or Conditions Relate
CURRENT APPLICATION NUMBER: US/09/685,664B
CURRENT APPLICATION NUMBER: US/09/685,664B
CURRENT FILING DATE: 1905-10-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-01-08
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR FILING DATE: 1995-01-08
PRIOR APPLICATION NUMBER: US 09/371,772
PRIOR APPLICATION NUMBER: US 09/371,772
PRIOR FILING DATE: 1999-08-10
NUMBER OF SEO ID NOS: 8231
SOFTWARE: Patentin version 3:0
LENGTH: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; PRIOR APPLICATION NUMBER: US 09/371,772
; PRIOR FILLING DATE: 1999-08-10
; NUMBER OF SEQ ID NOS: 8231
; SOFTWARE: PatentIn version 3.0
; SEQ ID NO 3109
; SEQ ID NO 3109
; LENGTH: 17
; TYPE: RNA
; ORGANISM: Mus musculus
US-09-685-664B-3109
                                                                                                                                                                                                                                                                                                                                       ; TYPE: RNA; Mus musculus US-09-685-664B-3110
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 15
US-09-685-664B-3110
Search completed: December 15, 2001, 05:58:20 Job time: 12263 sec
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                                                                                                                                                                                                               Query Match 100.0%; Score 7; DB 6; Length 17; Best Local Similarity 42.9%; Pred. No. 7.1e+04; Matches 3; Conservative 4; Mismatches 0; Indels
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7 uuugaua 13
                                                                                                                1 tttgata 7 :::||:| 3 uuugaua 9
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 2000000000
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Perfect score:
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Copyright (c) 1993 - 2000 Compugen Ltd.
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48.423 Million cell updates/sec
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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17	17	46	27	25	22	17	17	42	16	17	17	19	22	19	17	17	17	17	54	19	27	25	24	27	25	28	30	43	21	18	34	13	12	5	51	11	18	17	17	17	; ¤	,
-09-342-069A-38	-09-342-069-389	-60-130-180-313	-09-684-016-29	275	-09-553-094-2	-09-371-146A-9004	-09-304-517A-90	-60-096-409-134	-09-248-796	-09-362-510	-09-362-510-50840	-09-515-128	-09-577-409	-09-515-126	-09-362-510	2-510	-09-371-805	-09-371-805	-60-211-750	S-09-521-640-13137	-09-684-016-21639	-09-654-617-	-09-637-086A-2573	-09-684-016-18127	-09-654-617-18127	-09-705-926-	S-09-770-255-	-60-103-748-0	-09-540-213-	-09-411-	-60-013-529-	-08-978-620-	8-812-505-	-60-127-461-	S-60-184-813-	S-08-725-863-20	-09-497-191	S-09-380-826A-	26A-	-09-380-826A-	10	;
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APPLICANT: Chappel, Rod
TITLE OF INVENTION: LEPTOSPIRA PATHOGENS
FILE REFERENCE: DAVIE79.001APC
CURRENT APPLICATION NUMBER: US/09/380,826A
CURRENT FILING DATE: 199-11-22
PRIOR APPLICATION NUMBER: PCT/AU98/00145
PRIOR FILING DATE: 1998-03-06
PRIOR APPLICATION NUMBER: AU PO5494/97
PRIOR EILING DATE: 1997-03-07
NUMBER OF SEQ ID NOS: 26
SOFTWARR: FastSEQ for Windows Version 4.0
SEQ ID NO 6
LENGTH: 22
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US-09-380-826A-2
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US-09-380-826A-6
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                                                                    ; NAME/KEY: misc_feature

LOCATION: (1)...(22)

OTHER INFORMATION: n = A,T,C or C

OTHER INFORMATION: Primer for L.

US-09-380-825A-6
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PRIOR FILING DATE: 1997-03-07
NUMBER OF SEQ ID NOS: 26
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 2
LENGTH: 22
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                                                                                                                                                                                                                                                                                                                                                                               Sequence 6, Application US/09380826A GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Chappel, Rod
TITLE OF INVENTION: LEPTOSPIRA PATHOGENS
FILE REPERENCE: DAVLE79.001APC
CURRENT APPLICATION NUMBER: US/09/380,826A
CURRENT FILING DATE: 1999-11-22
PRIOR APPLICATION NUMBER: PCT/AU98/00145
PRIOR FILING DATE: 1998-03-06
          Query Match
Best Local Similarity
Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA
                                                                                                                                                            TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                              FEATURE:
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             63.6%; Score 14; DB 17; 100.0%; Pred. No. 7.3e+02;
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25
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US-09-304-517A-85089
US-09-654-617-291806
US-09-684-016-291806
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   Mismatches
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                              Length 22;
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RESULT 4
US-09-497-191-767/c
US-09-497-191-767/c
Sequence 767, Application US/09497191
Sequence 767, Application US/09497191
GENERAL INFORMATION:
APPLICANT: CHEN, Xianfeng
APPLICANT: SUBRAMANIAN, Gopalakrishnan
APPLICANT: ZIENG, Liansheng
TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS 84-89, 91-99
FILE REFERENCE: 2750-694P
CURRENT FILING DATE: 2000-02-03
UMBER OF SEQ ID NO.567
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO.767
LENGTH: 147
TYPE: NUMBER
TITLE PATENTIN VER. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: PCT/AU98/00145
PRIOR FILING DATE: 1998-03-06
PRIOR APPLICATION NUMBER: AU PO5494/97
PRIOR FILING DATE: 1997-03-07
NUMBER OF SEQ ID NOS: 26
SOCTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 7
LENGTH: 22
TYPE: DNA
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                                                                                                                                                ) OTHER INFORMATION: any n or Xaa = unknown US-09-497-191-767
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Best Local Similarity 63.0
Matches 14; Conservative
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TITLE OF INVENTION: LEPTOSPIRA PATHOGENS
FILE REFERENCE: DAVIE79.001APC
CURRENT APPLICATION NUMBER: US/09/380,826A
CURRENT FILING DATE: 1999-11-22
                                                              Query Match
Best Local Similarity
Matches 14; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Artificial Sequence
                                                                                                                                                                                                  ORGANISM: Arabidopsis thaliana
                                                                                                                                                                                  FEATURE:
                                                                                                                                                                                                                      TYPE: DNA
55 TGTTGGACTTGATCTTTTGATA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 tgttggatcacaagatttgata
                 1 tgttggannnnnnnntttgata
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                                                                  Conservative
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Pred. No. 7.
                                   22
   34
                                                                                   Score 14; DB 18;
Pred. No. 1.1e+03;
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                                                                     Mismatches
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                                                                                                     DB 18;
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US-08-725-863-2005; Sequence 2005, Application US/08725863

RESULT

GENERAL INFORMATION: APPLICANT:
APPLICANT:

Lane, John C. Delegeane, Angelo M. Stuart, Susan G. ...
Gooding, Douglas H.

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Best Local Similarity
Matches 14; Conserv
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                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
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ATTORNEY/AGENT INFORMATION:
NAME: CERRONE, MICHAEL C., Ph.D.
REGISTRATION NUMBER: 39,132
REFERENCE/DOCKET NUMBER: PD-0043 US
TELECOMMUNICATION INFORMATION:
TELECHONE: (415) 845-4166
TELECHONE: (415) 845-4166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (415) 845-4166
INFORMATION FOR SEQ ID NO: :
SEQUENCE CHARACTERISTICS:
LENGTH: 243 base pairs
TYPE: nucleic acid
CURRENT APPLICATION NUMBER: US/60/184,813
CURRENT FILING DATE: 2000-02-24
                                    APPLICANT: Hillman, Jennifer L. Fong, Willy Tuen TITLE OF INVENTION: NUCLEIC ACID SYNTHESIS AND MODIFICATION ENZYMES FILE REFERENCE: PT-0121 P
                                                                                     APPLICANT: Rosen, Bruce
APPLICANT: Shah, Purvi
APPLICANT: Chalup, Michael S.
                                                                                                                             APPLICANT:
                                                                                                                                          APPLICANT: Banville, Steve C.
APPLICANT: Bratcher, Shawn R.
APPLICANT: Dufour, Gerard E.
APPLICANT: Cohen, Howard J.
                                                                                                                                                                                                                                       APPLICANT:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Word Perfect 6.1 for Windows/MS-DOS 6.2
CURRENT APPLICATION DATA:
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APPLICANT: Delegeane, Ang
APPLICANT: Snable, James
TITLE OF INVENTION: POLYN
TITLE OF INVENTION: HUMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       IMMEDIATE SOURCE:
CLONE: 342263
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                       152 TGTTGGAATTTGCTGTTTGATA 173
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY:
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                                                                                                                                                                                                                                                                                                             200, Application US/60184813
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                                                                                                                                                                                                                    Russo, Frank D.
Spiro, Peter A.
                                                                                                                                                                                                                                                                      Hodgson, David M.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            63.6%;
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Pred. No.
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Roseberry, Ann M.
Wright, Rachel J.
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                                                                                                                         Yap, Pierre E.
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                                                                                                                                                                                                                                                          Jimmy Y.
                                                                                                        Stockdreher, Theresa
                                                                                                                                            Chen, Wensheng
Liu, Tommy
                                                                                                                                                                                                                                     Greenawalt, Lila B.
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                                                                                         Stefan
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; FEATURE:
; NAME/KEY: unsure
; LOCATION: 130
; OTHER INFORMATION: a
US-60-184-813-200
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                                                                                                                                                                                                                                                                                 RESULT
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CURRENT FILING DATE: 1999-04-01
NUMBER OF SEQ ID NOS: 8247
SOFTWARE: PERL Program
SEQ ID NO 5765
                                                                                                                                                                                                                                    Sequence 2536, Application US/08812505 GENERAL INFORMATION:
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Best Local Similarity
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LENGTH: 243
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APPLICANT: Thornton, Michael
TITLE OF INVENTION: NUCLEOTIDE SEQUENCES OF CANDIDA ALBICANS
FILE REFERENCE: PM-0007-2 P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PERL Program
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
ORGANISM: Homo sapiens
FEATURE: -
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APPLICANT:
APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 256
                             CORRESPONDENCE ADDRESS:
ADDRESSEE: INCYTE PHARMACEUTICALS, INC
                                                                                                                                       APPLICANT:
                                                                             TITLE OF INVENTION:
                                                                                                         APPLICANT:
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                                                             NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              152 tgttggaatttgctgtttgata 173
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                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
nes 14; Conserv
                 STREET:
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PALO ALTO
                                                                                                     Ito, Laura Y.
Akerblom, Ingrid E.
Akerblom, Ingrid E.
Delegeane, Angelo M.
Naughton, Rebecca E.
Klingler, Tod M.
                 3174 PORTER DRIVE
                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                   Gooding, Douglas H.
                                                                                                                                                                                     Stuart, Susan G.
                                                                                                                                                                                                  Stuve, Laura L.
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                                                                            POLYNUCLEOTIDES AND POLYPEPTIDES DERIVED FROM ASTROCYTOMA
                                                                                                                                                                                                                                                                                                                                                                                                     63.6%;
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Pred. No. 1
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Pred. No. 1.2e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                       DB 45;
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                                                                                                                                                                                                                                                                                                                                                                                                          .3e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                     Length 256;
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                                                                                                                                                                                                                                                                                                                                                                                          0;
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COMPUTER READABLE FORM:

94304

STATE: (

USA

CALIFORNIA

MEDIUM TYPE: Floppy disk

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                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 2536, Applic
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE: (415) 855-05:
TELEFAX: (415) 852-0195
INFORMATION FOR SEQ ID NO: :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 259 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       IMMEDIATE SOURCE:
CLONE: 865159
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REGISTRATION NUMBER: 33954
REFERENCE/DOCKET NUMBER: PD-
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 855-0555
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION: NAME: LUTHER, BARBARA J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA: APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION DATA:
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OPERATING SYSTEM:
SOFTWARE: Word Pe
                           COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Word Perfect 6.1 for Windows/MS-DOS 6.2
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: 1:
                                                                                                              COMPUTER READABLE FORM:
                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                              TITLE OF INVENTION: POLYNUCLEOTIDES AND POLYPEPTIDES DERIVED FROM
                                                                                                                                                                                                                                                                                              APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION:
                                                                                                                                                                                             STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          32 TGTTGGAATTTGTCATTTGATA 53
APPLICATION NUMBER: FILING DATE: HEREW
                                                                                                                                                                  STATE:
                                                                                                                                                                               CITY: PALO ALTO
                                                                                                                                             COUNTRY:
                                                                                                                                                                                                                ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 tgttggannnnnnntttgata 22
                                                                                            MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       14; Conservative
                                                                                                                               94304
                                                                                                                                                                                                                                                INVENTION: HUMAN BRAIN F SEQUENCES: 7797
                                                                                                                                                              CALIFORNIA
                                                                                                                                                                                               3174 PORTER DRIVE
                                                                                                                                               USA
                                                                                                                                                                                                                                                                                           Delegeane, Angelo M.
Naughton, Rebecca E.
Klingler, Tod M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Word Perfect 6.1 for Windows/MS-DOS 6.2
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SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                              Gooding, Dougl
Stuve, Laura L
                                                                                                                                                                                                                                                                                                                                                                                                                                          Application US/08978620
                                                                                                                                                                                                                                                                                                                                              Akerblom, Ingrid E.
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                                                                                                                                                                                                                INCYTE PHARMACEUTICALS, INC
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                                                                                               Floppy disk
                                                                                                                                                                                                                                                                                                                                                             Laura Y.
 HEREWITH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       63.68;
                                                                                                                                                                                                                                                                                                                                                                                                           Douglas H.
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               US/08/978,620
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEPHONE: (650) 855-0555
TELEFAX: (650) 852-0195
INFORMATION FOR SEQ ID NO: 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: LUTHER, BARBARA J
REGISTATION NUMBER: 33954
REFERENCE/DOCKET NUMBER: PD
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IMMEDIATE SOURCE: CLONE: 865159
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: 08/812
FILING DATE: MARCH 7, 1997
CLASSIFICATION: 536
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION: 536 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION: 536 PRIOR APPLICATION DATA:
                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Word Perfect 6.1 for Windows/MS-DOS 6.2
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
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               ATTORNEY/AGENT INFORMATION:
NAME: LUTHER, BARBARA J.
REGISTRATION UMBER: 33954
REFERENCE/DOCKET NUMBER: PI
                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES:
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FELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                     STREET: 3174 PORTE CITY: PALO ALTO STATE: CALIFORNIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local
                                                                           FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                          COUNTRY: UZIP: 94304
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             32 TGTTGGAATTTGTCATTTGATA 53
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                                                                                                             APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                       ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ENGTH:
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14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                 Akerblom, Ingrid E.
Delegeane, Angelo M.
Naughton, Rebecca E.
Klingler, Tod M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 259 base pairs
                                                                                                                                                                                                                                                                                          3174 PORTER DRIVE
                                                                                                                                                                                                                                           USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ito,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gooding, Douglas H. Stuve, Laura L. Stuart, Susan G.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   Laura
                                                                                                                                                                                                                                                                                                                                                   POLYNUCLEOTIDES AND POLYPEPTIDES DERIVED ASTROCYTOMA
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                                                                                                                                                                                                                                                                                                                                         3896
                                                                                                              us/60/013,529
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Pred. No.
                PD-0126P
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2536:

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APPLICANT: Stuve, Laura
APPLICANT: Goralski, Thomas J.
APPLICANT: Goralski, Thomas J.
FILE OF INVENTION: POLYNUCLEOTIDES AND POLYPEPTIDES DERIVED FROM RAT LIVER
FILE REFERENCE: PZ-0096 US
CURRENT APPLICATION NUMBER: US/09/411,077
CURRENT FILING DATE: 1999-10-04
EARLIER APPLICATION NUMBER: 60/103,748
EARLIER FILING DATE: October 9, 1998
EARLIER FILING DATE: October 9, 1998
EARLIER FILING DATE: December 10, 1998
EARLIER APPLICATION NUMBER: 60/111,900
EARLIER FILING DATE: December 10, 1998
INUMBER OF SEQ ID NOS: 1651
SOFTWARE: PERL Program
SEQ ID NO 732
LENGTH: 261
TYPE: DNA
ORGANISM: Rattus norvegicus
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: OTHER INFORMATION: 701257277H1
US-09-411-077-732
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                                                                                                                                                                                                                                                   US-09-540-213-52141; Sequence 52141, Application US/09540213
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-09-411-077-732
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TELEFAX: (415) 852-0195
INFORMATION FOR SEQ ID NO: 2536
SEQUENCE CHARACTERISTICS:
LENGTH: 259 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                   GENERAL INFORMATION:
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Best Local Similarity
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Best Local Similarity
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                              APPLICANT: Mullahy, Sara J.
APPLICANT: Naughton, Rebecca E.
TITLE OF INVENTION: POLYNUCLEOTIDES OF LIVER
FILE REFERENCE: PD-1031 CIP
CURRENT APPLICATION NUMBER: US/09/540,213
CURRENT FILING DATE: 2000-03-31
                                                                                                                                                    APPLICANT: Seilhamer, Jeffrey J.
APPLICANT: DeLegeane, Angelo M.
APPLICANT: Stuart, Susan G.
APPLICANT: Stuve, Laura L.
Prior application data removed - consult PALM or file wrapper NUMBER OF SEQ ID NOS: 53844
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                                                                                                                                                                                                                                                                                                                                                                                                                              14; Conservative
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Pred. No. 1.3e+03;
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Pred. No. 1.3e+03
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, NAME/KEY: misc_feature
; OTHER INFORMATION: Incyte ID No: rat00178536
US-09-540-213-52141
                                                                                                                                                                                                                                                                                                       RESULT 14
US-09-770-255-141
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Rattus norvegicus; FEATURE: -
OTHER INFORMATION: 701257277H1
US-60-103-748-618
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APPLICANT: Stuve, Laura L.
APPLICANT: Goralski, Thomas J.
APPLICANT: Garrow, Bonnie L.
TITLE OF INVENTION: POLYNUCLEOTIDES AND POLYPEPTIDES DERIVED FROM TREATED RAT LIVE
FILE REFERENCE: PZ-0096 P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/60/103,748
CURRENT FILING DATE: 1998-10-09
NUMBER OF SEQ ID NOS: 1537
SOFTWARE: PERL Program
SEQ ID NO 618
LENGTH: 261
TYPE: DNA
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Best Local Similarity
Matches 14; Conserv
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Best Local :
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TYPE: DNA
ORGANISM: Rattus norvegicus
                              APPLICANT
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               APPLICANT
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                                                                                               Matthew, Abraham V.
Ledford, Brooke L.
Woessner, Jeffrey P.
Haas, William David
                                                                                                                                                                                                                                                              Gorlach, Jorn
                                                     Slader,
                                                                                                                                                                                                                                 An, Yong-Qiang
Hamilton, Carol
Hurban, Patrick
                            Davis, Keith R. Allen, Keith
                                                        Kricker, Maja
Slader, Ted
                                                                                     Garcia, Carlos A.
                                                                                                                                                           Page, Amy
                                                                                                                                                                      Yu, Yang
Rameaka, Joshua G.
                                                                                                                                                                                                      Raines,
                                                                                                                                                                                                                   Price, Jennifer L.
             Hoi fman,
                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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                                                                                                                                                                                                   Tracy M.
                                                                                                                                                                                                                                                                                                                                                                                                                                         63.68;
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Pred. No. 1.
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TITLE OF INVENTION: Expressed Sequences of Arabidopsis
FILE REFERENCE: 2030US (PARA-019PRV)
CURRENT APPLICATION NUMBER: US/09/770,255
CURRENT FILING DATE: 2001-01-26
PRIOR APPLICATION NUMBER: 60/178.504
PRIOR PELLING DATE: 2000-01-27
NUMBER OF SEQ ID NOS: 999
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 141
LENGTH: 317
TYPE: DNA
ORGANISM: Arabidopsis thaliana
FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)...(317)
OTHER INFORMATION: n = A,T,C or G
US-09-770-255-141
                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:

APPLICANT: LIU, Jingdong

APPLICANT: Wiegand, Roger C.

APPLICANT: Wiegand, Roger C.

TITLE OF INVENTION:

FILE REFERENCE: 38-21(12481)A

CURRENT APPLICATION NUMBER: US/09/705,926

CURRENT FILING DATE: 2000-11-06

NUMBER OF SEQ ID NOS: 21634

SEQ ID NO 17256

LENGTH: 322

TYPE: DNA

ORGANISM: Lycopersicon esculentum
FEATURE:

NAME/KEY: Unsure

LOCATION: (1)..(322)

OTHER INFORMATION: unsure at all n locations
US-09-705-926-17256
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 15
US-09-705-926-17256/c
Sequence 17256, Application US/09705926
GENERAL INFORMATION:
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Search completed: December 15, 2001, 05:42:59 Job time: 15824 sec
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Best Local Similarity 63.6%;
Matches 14; Conservative
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                                                                                                      1 tgttggannnnnnntttgata 22
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|12 TGTTGGAAGGGTTCTTTTGATA 91
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                                                                                                                                                                                             Score 14; DB 28; Length 322; pred. No. 1.3e+03; 0; Mismatches 8; Indels
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Result
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Maximum Match 100%
Listing first 45 summaries
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   Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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seq length: 2000000000
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Match
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                      Pending_Patents_NA_New: *
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Gapop 10.0 , Gapext 1.0
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22
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Copyright (c) 1993 - 2000 Comp
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/cgn2_6/ptodata/1/pna/US06_NEW_COMB.seq:*
/cgn2_6/ptodata/1/pna/US07_NEW_COMB.seq:*
/cgn2_6/ptodata/1/pna/US09_NEW_COMB.seq:*
/cgn2_6/ptodata/1/pna/US09_NEW_COMB.seq:*
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6 US-09-985-678-90042
6 US-09-985-678-85089
6 US-09-922-279A-3008
1 PCT-US01-15674A-50
6 US-09-815-264-12871
5 US-09-815-264-50077
7 US-09-815-264-60943
8 US-09-815-264-74657
8 US-09-815-264-74657
8 US-09-815-264-79220
9 US-09-815-264-74657
9 US-09-815-267-74151011
9 US-09-815-242-7111
9 US-09-815-242-7111
9 US-09-933-524A-18588
9 US-09-933-524A-18588
9 US-09-933-524A-18588
9 US-09-933-524A-110244
9 US-09-815-264-29506
9 US-09-815-264-29506
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                              1500, Ap
3155, Ap
2111, Ap
231502,
23742, A
18588, A
18588, App
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12871, Appl

12871, A

50027, A

50028, A

50043, A

60943, A

69885, A

79259, A

79259, A

11444, Ap
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151011,
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ALIGNMENTS	US-09-815-264-78579	US-09-815-264-66587	US-09-815-264-58271	US-09-815-264-91108	US-09-815-242-6647	US-09-897-516-970	US-09-815-264-106280	US-09-815-264-25196	US-09-815-264-7540	US-09-815-264-106283	US-09-898-888A-14137	US-09-815-242-8747	US-09-815-242-8090	US-09-815-242-4274	US-09-922-279A-3103	US-09-815-242-4840	US-09-922-279A-1268	US-09-815-264-65443	US-09-985-678-220277	US-09-933-524A-109421
	Sequence 78579, A	Sequence 66587, A	Sequence 58271, A	Sequence 91108, A	Sequence 6647, Ap	Sequence 970, App	Sequence 106280,	Sequence 25196, A	Sequence 7540, Ap	Sequence 106283,	Sequence 14137, A	Sequence 8747, Ap	Sequence 8090, Ap	Sequence 4274, Ap			Sequence 1268, Ap	Sequence 65443, A	Sequence 220277,	Sequence 109421,

POTOMITEMA

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US-09-985-678-90042/c

Sequence 90042, Application US/09985678

GENERAL INFORMATION:
APPLICANT: Cheikh, Nordine
APPLICANT: Liu, Jingdong
TITLE OF INVENTION: Annotated Plant Genes
FILE REFERENCE: 16517.255,38-21(15097)F
CURRENT APPLICATION NUMBER: US/09/985,678
CURRENT APPLICATION NUMBER: US/09/985,678
CURRENT FILING DATE: 2001-11-05
PRIOR APPLICATION NUMBER: US 09/304,517
PRIOR FILING DATE: 1999-05-06
NUMBER OF SEQ ID NOS: 295529
SEQ ID NO 90042
LENGTH: 427
APPLICANT: Liu, Jingdong
TITLE OF INVENTION: Annotated Plant Genes
FILE REFERENCE: 16517.255/38-21(15097)F
CURRENT APPLICATION NUMBER: US/09/985.678
CURRENT FILING DATE: 2001-11-05
PRIOR APPLICATION NUMBER: US 09/304,517
PRIOR FILING DATE: 1999-05-06
NUMBER OF SEQ ID NOS: 295529
SEQ ID NO 85089
LENGTH: 461
TYPE: DNA
ORGANISM: Zea mays
US-09-985-678-85089
                                                                                                                                                                                                                                                                                     RESULT 2
US-09-985-678-85089/c
Sequence 85089, Application US/09985678
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; TYPE: DNA
; ORGANISM: Zea mays
US-09-985-678-90042
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Best Local :
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nes 14; Conserv
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Pred. No. 2
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2.2e+02;
8;
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                                                                                                                                                                                                                                                                                                                                             PCT-US01-15674A-50
                                                                                       Sequence 50, Application PC/TUS0115674A
GENERAL INFORMATION:
APPLICANT: JOAN, KNOLL H
APPLICANT: ROGAN, PETER K
TITLE OF INVENTION: SINGLE COPY GENOMIC HYBRIDIZATION PROBES AND METHOD OF GENERATING FILE REFERENCE: 30307
CURRENT APPLICATION NUMBER: PCT/US01/15674A
CURRENT FILING DATE: 2001-12-07
NUMBER OF SEQ ID NOS: 613
SOFTWARE: Patentin version 3.1
SEQ ID NO 50
LENGTH: 1287
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/09/922,279A CURRENT FILING DATE: 2001-08-03 PRIOR APPLICATION NUMBER: 09/491,404 PRIOR FILING DATE: 2000-01-25 NUMBER OF SED ID NOS: 3796 SOFTWARE: pt_SP_genes Version 1.0 SEQ ID NO 3008 LENGTH: 857
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity Matches 14; Conserv
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APPLICANT: Drmanac, Radoje T.
TITLE OF INVENTION: Novel Contigs Obtained
TITLE OF INVENTION: From Various Libraries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Tang, Yuanhua T.
APPLICANT: Tillinghast, John
APPLICANT: Sinku, Ankura
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: sig_peptide
LOCATION: (514)...(649)
OTHER INFORMATION: this location contains the signal peptide sequence,
OTHER INFORMATION: MGAFVTRVALPCSFAWCACILGPASALCSLCMKNFQPFFQVILDS, Run with SignalP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILE REFERENCE: 785
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OTHER INFORMATION: similar to gi2231098 in the genepept database release 114 OTHER INFORMATION: Run with FASTXY 3.3t00, default parameters
NAME/KEY: repeat_region LOCATION: (1)..(1287) OTHER INFORMATION: hsmar1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: misc_feature LOCATION: (290)...(213)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
ORGANISM: Homo sapiens
                                                       FEATURE:
                                                                     ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                     685 tgttggaggaagtgctttgata
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 14;
Pred. No.
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PRIOR APPLICATION NUMBER: US 09/620,392
PRIOR FILING DATE: 2000-07-19
PRIOR APPLICATION NUMBER: US 09/702,134
PRIOR FILING DATE: 2000-10-31
NUMBER OF SEQ ID NOS: 109669
SEQ ID NO 12871
LENGTH: 1501
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-09-815-264-12871; Sequence 12871, Application US/09815264; GENERAL INFORMATION:
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US-09-815-264-12871
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    Matches

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Best Local Similarity
Matches 14; Conserv
          NUMBER OF SEQ ID NOS: 109669
SEQ ID NO 14471
LENGTH: 1501
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                                                                                        APPLICANT: Wu, Wei
TITLE OF INVENTION: Genomic Plant Sequences
FILE REFERENCE: 38-21(51237)G
CURRENT APPLICATION NUMBER: US/09/815,264
CURRENT FILING DATE: 2001-03-23
PRIOR APPLICATION NUMBER: US 09/620,392
PRIOR FILING DATE: 2000-07-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Wu, Wei
ITILE OF INVENTION: Genomic Plant Sequences And Uses Thereof
FILE REFERENCE: 38-21(51237)G
CURRENT APPLICATION NUMBER: US/09/815,264
CURRENT FILING DATE: 2001-03-23
                                                          PRIOR APPLICATION NUMBER: US 09/702,134 PRIOR FILING DATE: 2000-10-31
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                                                                                                                                                                                                                                                                                                                                                                                                               671 tgttggaacaggaattttgata
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                   1 tgttggannnnnnnntttgata 22
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DNA
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Koshi, Jeffrey M.
Kovalic, David K.
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Dotson, Stanton B.
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Pred. No. 2.6e+02;
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2.5e+02;
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; ORGANISM: Oryza sativa US-09-815-264-14471

Query Match 63.6%; Best Local Similarity 63.6%;

Score 14; DB 6; Pred. No. 2.6e+02;

Length 1501;

0;

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APPLICANT: WGININGH, James

APPLICANT: WU, Wei

APPLICANT: WWI, Wei

FILE REFERENCE: 38-21(51237)G

CURRENT APPLICATION NUMBER* US/09/815,264

CURRENT FILING DATE: 2001-03-23

PRIOR APPLICATION NUMBER: US 09/620,392

PRIOR FILING DATE: 2000-07-19

PRIOR FILING DATE: 2000-10-31

PRIOR FILING DATE: 2000-10-31

PRIOR FILING DATE: 2000-10-31

NUMBER OF SEO ID NOS: 109669
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PRIOR FILING DATE: 2000-07-19
PRIOR APPLICATION NUMBER: US 09/702,134
PRIOR FILING DATE: 2000-10-31
NUMBER OF SEO ID NOS: 109669
SEO ID NO 50077
LENGTH: 1501
TYPE: DNA
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US-09-815-264-50077
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GENERAL INFORMATION
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Best Local Similarity
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APPLICANT: Cao, Yongwei
APPLICANT: Dotson, Stanton B.
APPLICANT: Koshi, Jeffrey M.
APPLICANT: Kovalio, David K.
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Koshi, Jeffrey M.
Kovalic, David K.
Liu, Jingdong
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Koshi, Jeffrey M.
Kovalic, David K.
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Pred. No. 2.6e+02;
Pred. No. 2.6e+02;
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TITLE OF INVENTION: Genomic Plant Sequences And Uses Thereof
FILE REFERENCE: 38-21(51237)G
CURRENT APPLICATION NUMBER: US/09/815,264
CURRENT FILING DATE: 2001-03-23
PRIOR APPLICATION NUMBER: US 09/620,392
PRIOR APPLICATION NUMBER: US 09/620,392
PRIOR APPLICATION NUMBER: US 09/702,134
PRIOR FILING DATE: 2000-07-19
PRIOR APPLICATION NUMBER: US 09/702,134
PRIOR FILING DATE: 2000-10-31
NUMBER OF SEQ ID NOS: 109669
SEQ ID NO 60943
LENGTH: 6926
TYPE: DNA
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US-09-815-264-50328
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US-09-815-264-60943
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APPLICANT: BOUKharcov, Andrey A.
APPLICANT: Cao, Yongwei
APPLICANT: Dotson, Stanton B.
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APPLICANT:
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Best Local Similarity
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SEQ ID NO 69885
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             PRIOR APPLICATION NUMBER: US 09/702,134
PRIOR FILING DATE: 2000-10-31
NUMBER OF SEQ ID NOS: 109669
                                                                                                                          APPLICANT: McIninch, James
APPLICANT: Wu, Wei
TITLE OF INVENTION: Genomic Plant Sequences
FILE REFERENCE: 38-21(51237)/6
FILE REFERENCE: 38-21(51237)/6
FILE REFERENCE: 38-21(51237)/6
                                                                    CURRENT APPLICATION NUMBER: US/09/815,264 CURRENT FILING DATE: 2001-03-23 PRIOR APPLICATION NUMBER: US 09/620,392 PRIOR FILING DATE: 2000-07-19
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Kovalic, David K.
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Pred. No. 2.6e+02;
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у.2e+02;
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US-09-815-264-74657/c
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US-09-815-264-69885
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US-09-815-264-74657
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TITLE OF INVENTION: Genomic Plant Sequences An
FILE REFERENCE: 38-21(51237)G
CURRENT APPLICATION NUMBER: US/09/815,264
CURRENT FILING DATE: 2001-03-23
PRIOR APPLICATION NUMBER: US 09/620,392
PRIOR FILING DATE: 2000-07-19
PRIOR APPLICATION NUMBER: US 09/702,134
PRIOR FILING DATE: 2000-10-31
PRIOR FILING DATE: 2000-10-31
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Best Local Similarity 63.6
Matches 14; Conservative
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SEQ ID NO 74657
LENGTH: 7673
TYPE: DNA
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Best Local
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          TITLE OF INVENTION: Genomic Plant Sequences and Uses Thereof FILE REFERENCE: 38-21(51237)6
CURRENT APPLICATION NUMBER: US/09/815;264
CURRENT FILING DATE: 2001-03-23
PRIOR APPLICATION NUMBER: US 09/620,392
PRIOR PILING DATE: 2000-07-19
PRIOR FILING DATE: 2000-07-19
PRIOR FILING DATE: 2000-10-31
PRIOR FILING DATE: 2000-10-31
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               14;
SEQ ID NOS:
                                                                                                                                                                Koshi, Jeffrey M.
Kovalic, David K.
Liu, Jingdong
McIninch, James
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Koshi, Jeffrey M.
Kovalic, David K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Dotson,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      McIninch, James
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Liu, Jingdong
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Application US/09815264
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Stanton B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      63.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                63.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 14;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 6;
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; NAME/KEY: unsure
; LOCATION: (1)..(41742)
; OTHER INFORMATION: unsure at all n locations
US-09-815-264-59220
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Best Local Similarity
Thes 14; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; ORGANISM: Oryza sativa 
US-09-815-264-79259
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-09-815-264-79259
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT
                                                                                                                                                                                                                                  US-09-815-242-1444
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 79259, Application US/09815264 GENERAL INFORMATION:
                                                                                                                                                                                               Sequence 1444, Application US/09815242 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ ID NO 79259
LENGTH: 94612
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQ ID NO 59220
LENGTH: 41742
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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Best Local Similarity
                                                                            APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Wu, Wei
TITLE OF INVENTION: Genomic Plant Sequences
FILE REFERENCE: 38-21(51237)G
CURRENT APPLICATION NUMBER: US/09/815,264
CURRENT FILING DATE: 2001-03-23
PRIOR APPLICATION NUMBER: US 09/620,392
PRIOR FILING DATE: 2000-07-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Boukharov, Andrey A. APPLICANT: Cao, Yongwei
                                                                                                                                                APPLICANT: Haselbeck, Robert APPLICANT: Ohlsen, Kari L. APPLICANT: Zyskind, Judith V
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: US 09/702,134 PRIOR FILING DATE: 2000-10-31
               TITLE OF INVENTION: Identification of Essential Genes TITLE OF INVENTION: Prokaryotes FILE REFERENCE: ELITRA,011A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQ ID NOS: 109669
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
CURRENT APPLICATION NUMBER: US/09/815,242
                                                                APPLICANT:
                                                                                                                              APPLICANT:
                                                                                                                                                                                                                                                                                                    14796 tgttggatattatcctttgata 14817
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Oryza sativa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
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Koshi, Jeffrey M.
Kovalic, David K.
Liu, Jingdong
                                                                            Wall, Daniel
Trawick, John D.
Carr, Grant J.
Yamamoto, Robert T.
                                                                  Xu, H. Howard
                                                                                                                                                Ohlsen, Kari L.
Zyskind, Judith W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cao, Yongwei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 McIninch, James
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 14; DB 6;
Pred. No. 4.1e+02;
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CURRENT FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: 60/191,078
PRIOR FILING DATE: 2000-03-21
PRIOR APPLICATION NUMBER: 60/206,848
PRIOR FILING DATE: 2000-05-23
PRIOR APPLICATION NUMBER: 60/207,727
PRIOR APPLICATION NUMBER: 60/242,578
PRIOR APPLICATION NUMBER: 60/242,578
PRIOR FILING DATE: 2000-10-23
PRIOR APPLICATION NUMBER: 60/253,625
PRIOR FILING DATE: 2000-11-27
PRIOR APPLICATION NUMBER: 60/257,931
PRIOR FILING DATE: 2000-12-22
PRIOR FILING DATE: 2000-12-22
PRIOR APPLICATION NUMBER: 60/269,308
PRIOR FILING DATE: 2001-02-16
NUMBER OF SEQ ID NOS: 14110
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1444
LENGTH: 226
TYPE: DNA
ORGANISM: Staphylococcus aureus
US-09-815-242-1444
                                                                                                                                                                                                                                                                                                                                                 RESULT 15
US-09-985-678-151011/c
Sequence 151011, Application US/09985678
GENERAL INFORMATION:
APPLICANT: Cheikh, Nordine
APPLICANT: Liu, Jingdong
TITLE OF INVENTION: Annotated Plant Genes
FILE REFERENCE: 16517.255/38-21(15097)F
CURRENT APPLICATION NUMBER: US/09/985,678
CURRENT FILING DATE: 1909-05-06
PRIOR APPLICATION NUMBER: US 09/304,517
PRIOR FILING DATE: 1999-05-06
NUMBER OF SEQ ID NOS: 295529
SEQ ID NO 151011
LENGTH: 245
TYPE: DNA
ORGANISM: Glycine max
US-09-985-678-151011
Search completed: December 15, 2001, 05:58:24 Job time: 12267 sec
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ouery Match 59.1%; Score 13; DB 6; Length 226 Best Local Similarity 61.9%; Pred. No. 6.9e+02; Matches 13; Conservative 0; Mismatches 8; Indels
                                                                                                                       2 gttggannnnnnntttgata 22
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165 GTTGGATTGTATCATTTGATA 145
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Minimum DB seq
Maximum DB seq
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Perfect score:
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Maximum Match 100%
Listing first 45 s
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1: /cgn2_6/ptodata/2/pna/US

2: /cgn2_6/ptodata/2/pna/US

3: /cgn2_6/ptodata/2/pna/US

4: /cgn2_6/ptodata/2/pna/US

5: /cgn2_6/ptodata/2/pna/US

6: /cgn2_6/ptodata/2/pna/US

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11: /cgn2_6/ptodata/2/pna/US
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length: 2000000000
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Gapop 10.0 , Gapext 1.
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cgn2_6/ptodata/2/pna/US08_COMB.seq: *
cgn2_6/ptodata/2/pna/US09_COMB.seq: *
cgn2_6/ptodata/2/pna/US09_COMB.seq: *
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cgn2_6/ptodata/2/pna/US0006_COMB.seq: *
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/cgn2_6/ptodata/2/pna/US06_COMB.seq:*
/cgn2_6/ptodata/2/pna/US07_COMB.seq:*
/cgn2_6/ptodata/2/pna/US080_COMB.seq:*
/cgn2_6/ptodata/2/pna/US081_COMB.seq:*
/cgn2_6/ptodata/2/pna/US082_COMB.seq:*
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SUMMARIES

Query

Result No. ဂဂ O ဂ ဂ O Score Match Length DB 7 US-09-684-016-175385 2 US-09-565-309A-57118 2 US-09-565-309A-57118 2 US-09-565-309A-57118 2 US-09-684-016-117747 5 US-09-684-016-117747 6 US-09-404-520-528 8 US-09-404-520-528 8 US-09-983-36-352 1 US-09-983-859-3148 2 US-60-1191-637-27704 2 US-60-1191-637-27704 2 US-60-1191-637-217-19072 2 US-60-1191-637-217-19072 2 US-60-1191-637-217-18350 3 US-60-1191-637-11908 7 US-60-1191-637-118351 2 US-60-1191-637-118350 3 US-60-1191-637-118351 3 US-09-684-617-1180482 9 US-09-684-617-1180482 9 US-09-684-617-1180482 9 US-09-684-617-1180482 9 US-09-534-859-802 1 US-09-803-736-23 ij US-09-380-826A-1 Sequence 1, Appli Sequence 11944, A Sequence 175385, Sequence 20908, A Sequence 20908, A Sequence 57118, A Sequence 117747, Sequence 11852, App Sequence 1281, App Sequence 1650, App Sequence 19008, A Sequence 19008, A Sequence 18350, App Sequence 18350, A Sequence 18350, A Sequence 18350, A Sequence 18350, A Sequence 180482, A Sequence 180482, A Sequence 103292, App Sequence 103292, Sequence 23, Appl Sequence 951, App Sequence 951, App Sequence 951, App Description Sequence Sequence

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Sequence 7, Application US/09380826A
GENERAL INFORMATION:
APPLICANT: Chappel, Rod
TITLE OF INVENTION: LEPTOSPIRA PATHOGENS
FILE REFERENCE: DAVIE79, 001APC
CURRENT APPLICATION NUMBER: US/09/380,826A
CURRENT FILING DATE: 1999-11-22
PRIOR APPLICATION NUMBER: PCT/AU98/00145
PRIOR FILING DATE: 1998-03-06
PRIOR FILING DATE: 1998-03-06
                                                                                                                                                               PRIOR APPLICATION NUMBER: AU PO5494/97
PRIOR FILING DATE: 1997-03-07
NUMBER OF SEQ ID NOS: 26
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 7
LENGTH: 22
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Chappel, Rod
TITILE OF INVENTION: LEPTOSPIRA PATHOGENS
FILE REFERENCE: DAVLET9.001APC
CURRENT APPLICATION NUMBER: US/09/380,826A
CURRENT FILING DATE: 1999-11-22
PRIOR APPLICATION NUMBER: PATHON PRIOR FILING DATE: 1998-03-06
PRIOR FILING DATE: 1998-03-06
PRIOR FILING DATE: 1997-03-07
PRIOR FILING DATE: 1997-03-07
NUMBER OF SEQ ID NOS: 26
SECTIMARE: FastSEQ for Windows Version 4.0
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Best Local Similarity
Watches 22; Conserva
                                                                                                                         , OTHER INFORMATION: Primer for L. US-09-380-826A-7
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LENGTH: 22
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
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                                                                     Query Match
Best Local Similarity
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                 1 tgttggatcacaagatttgata 22
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US-09-924-035-520
US-09-654-617-370318
US-09-684-016-370318
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Pred. No.
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Pred. No. 2;
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; TYPE: DNA; ORGANISM: Leptospira
US-09-380-826A-1
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US-09-380-826A-1
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; ORGANISM: Aspergillus nidulans
US-60-132-861-11944
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Cao, Yongwei
APPLICANT: Timberlake, William E.
APPLICANT: Timberlake, William E.
TITLE OF INVENTION: ASPERGILLUS NIDULANS GENOME SEQUENCE AND USES THEREOF
FILE REFERENCE: 38-21(15485)C
CURRENT APPLICATION NUMBER: US/60/132,861
CURRENT FILING DATE: 1999-05-07
NUMBER OF SEQ ID NOS: 13800
SEQ ID NO 11944
LENGTH: 420
                                                                                                                                                                                   Sequence 7425, Application US/09649165A GENERAL INFORMATION:
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LENGTH: 1
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Best Local Similarity
Matches 22; Conserv
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TITLE OF INVENTION: LEPTOSPIRA PATHOGENS
FILE REFERENCE: DAVIET9.001APC
CURRENT APPLICATION NUMBER: US/09/380,826A
CURRENT FILING DATE: 1999-11-22
PRIOR APPLICATION NUMBER: PCT/AU98/00145
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APPLICANT: Guegler, Karl J.
APPLICANT: Guegler, Martin
APPLICANT: Momiyama, Monika
APPLICANT: Momiyama, Monika
APPLICANT: Wang, Xinhao
TITLE OF INVENTION: DETECTING EXPRESSION OF ARABIDOPSIS THALIANA GENES
FILE REFERENCE: PA-0019 US
CURRENT APPLICATION NUMBER: US/09/649,165A
CURRENT FILING DATE: 2000-08-24
PRIOR APPLICATION NUMBER: 60/150,517
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Pred. No. 1
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1.1e+02;
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PRIOR FILING DATE: 1999-08-24

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PRIOR APPLICATION NUMBER:

PRIOR FILING DATE: 2000-09-
NUMBER OF SEQ ID NOS: 46317
SEQ ID NO 175385
LENGTH: 581
TYPE: DNA
ORGANISM: Arabidopsis thaliana
US-09-684-016-175385
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LENGTH: 581
: TYPE: DIA
: ORGANISM: Arabidopsis thaliana
US-09-654-617-175385
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: KOVE
APPLICANT: Liu, Jing
TITLE OF INVENTION:
FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
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SOFTWARE: PERL Program
SEQ ID NO 7425

LENGTH: 530

TYPE: DNA
ORGANISM: Arabidopsis thaliana
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Best Local Similarity
Matches 20; Conserv
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                                                                                                               FILE REFERENCE: 38-21(15097)D CURRENT APPLICATION NUMBER: US/09/684,016 CURRENT FILING DATE: 2000-10-10
                                                                                                                                            APPLICANT: Kovalic, APPLICANT: Liu, Jingdong TITLE OF INVENTION: FILE REFERENCE: 38-
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LOCATION: 6
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OTHER INFORMATION: Incyte ID No: 701676626
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2000-09-05
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                                                                                                                                                                                                                                                                                           441
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Pred. No. 1.1e+02;
0; Mismatches 2;
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Pred. No. 1.1e+02;
0; Mismatches 2;
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. 1.1e+02;
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Query Match

85.5%;

Score 18.8;

DΒ 27;

Length 581;

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US-09-565-309A-57118, Application US/09565309A; Sequence 57118, Application US/09565309A; GENERAL INFORMATION:
APPLICANT: ALEXANDROV, Nickolai; APPLICANT: BROVER, Vyacheslav
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Best Local Similarity
Thehes 20; Conserve
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. LCCATION: (1)..(649)
. OTHER INFORMATION: 27496:110925
US-09-565-309A-20908
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CURRENT FILING DATE: 2000-05-05
NUMBER OF SEQ ID NOS: 68449
SEQ ID NO 20908
LENGTH: 649
TYPE: DNA
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GENERAL INFORMATION:
APPLICANT: ALEXANDROV, Nickolai
APPLICANT: BROVER, Vyacheslav
                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/565,309A
CURRENT FILING DATE: 2000-05-05
NUMBER OF SEQ ID NOS: 68449
SEQ ID NO 57118
LENGTH: 1069
TYPE: DNA
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              Query Match
Best Local Similarity
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TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTID
TITLE OF INVENTION: THEREBY
FILE REFERENCE: 2750-0853P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: BROVER, VYACHESLAV
TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTID
TITLE OF INVENTION: THEREBY
FILE REFERENCE: 2750-0853P
                                                                               NAME/KEY: misc_feature
LOCATION: (1):.(1069)
OTHER INFORMATION: 27496 : OVERLAP (Clone Number : OVERLAP)
                                                                                                                                            FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)..(1069)
OTHER INFORMATION: any n =
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                                                                                                                                                                                                               ORGANISM: Arabidopsis thaliana
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0; Mismatches 2;
            Score 18.8;
Pred. No. 1.
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Pred. No. 1.
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NAME/KEY: misc_feature
COCATION: (1).(1072)
COTHER IMFORMATION: any n = a,
NAME/KEY: misc_feature
COCATION: (1).(1072)
OTHER IMFORMATION: 27496 : 51
US-09-565-309A-49885
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                                                                                                                                                                                                                            ; TYPE: DNA; ORANISM: Arabidopsis thaliana; OTHER INFORMATION: unsure at all n locations US-09-654-617-117747
                                                                                                                                                                                                                                                                                                                                                                                                                           US-09-654-617-117747; Application US/09654617; Sequence 117747, Application US/09654617
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                                         RESULT 12
US-09-684-016-117747
                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/54,617; CURRENT FILING DATE: 2000-09-05 NUMBER OF SEQ ID NOS: 463173
SEQ ID NO 117747
LENGTH: 1177
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LENGTH: 1072
              Sequence 117747, Ap GENERAL INFORMATION
                                                                                                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
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Best Local
                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: ALEXANDROV, Nickolai
APPLICANT: BROVER, Vyacheslav
TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTIDES
TITLE OF INVENTION: THEREBY
FILE REFERENCE: 2750-0853P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/565,309A
CURRENT FILING DATE: 2000-05-05
NUMBER OF SEQ ID NOS: 68449
                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Kovalic, APPLICANT: Liu, Jingdong
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Arabidopsis thaliana
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| 719 tgttggttcacaagatttgaca 740
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les 20; Conserv
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                             Application US/09684016
 Kovalic, David K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      90.9%;
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Pred. No. 1
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; TYPE: DNA
; CRGANISM: Arabidopsis thaliana
; FEATURE:
; NAME/KEY: unsure
; LOCATION: (1)..(1128)
; OTHER INFORMATION: unsure at all n locations
US-09-684-016-117747
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Best Local Similarity
Matches 20; Conserv
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SEQ ID NO 11564
LENGTH: 1429
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQ ID NO 117747
LENGTH: 1128
                   NUMBER OF SEQ ID NOS: 44345
SEQ ID NO 5289
LENGTH: 5874
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Best Local Similarity
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                                                                 APPLICANT: Cao, Yongwei
APPLICANT: Ghodssi, Azita
APPLICANT: Hinkle, Gregory J.
APPLICANT: McIninch, James
APPLICANT: Timberlake, William E.
APPLICANT: Yu, Jaehyuk
TITLE OF INVENTION: Emericella nidulans Genome Sequence i
FILE REFERENCE: 38-10 (15498)A
CURRENT FILING DATE: 1999-09-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Cao, Yongwei
APPLICANT: Timberlake, William E.
APPLICANT: Timberlake, William E.
TITLE OF INVENTION: ASPERGILLUS NIDULANS GENOME SEQUENCE AND USES THEREOF
FILE REFERENCE: 38-10(15485)D
CURRENT APPLICATION NUMBER: US/60/138,103
CURRENT FILING DATE: 1999-06-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: PRIOR FILING DATE: NUMBER OF SEQ ID NOS:
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TYPE: DNA
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Pred.
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Pred. No. 1.2e+02
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RESULT 15
US-09-534-859-352
; Sequence 352, Application US/09534859
; GENERAL INFORMATION:
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Search completed: December 15, 2001, 05:43:01 Job time: 15826 sec
                                                                               Ouery Match 85.5%; Score 18.8; DB 20; Best Local Similarity 90.9%; Pred. No. 2e+02; Matches 20; Conservative 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                             Ouery Match 85.5%; Score 18.8; DB 18; Best Local Similarity 90.9%; Pred. No. 1.4e+02; Matches 20; Conservative 0; Mismatches 2;
                                           Indels
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  is derived by analysis of the total score distribution.
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1: /cgn2_6/ptodata/1/pna/U

2: /cgn2_6/ptodata/1/pna/U

3: /cgn2_6/ptodata/1/pna/U

4: /cgn2_6/ptodata/1/pna/U

5: /cgn2_6/ptodata/1/pna/U

6: /cgn2_6/ptodata/1/pna/U

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Listing first 45 summaries
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Copyright (c) 1993 - 2000 Comp
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/cgn2_6/ptodata/1/pna/US06_NEW_COMB.seq:*
/cgn2_6/ptodata/1/pna/US08_NEW_COMB.seq:*
/cgn2_6/ptodata/1/pna/US08_NEW_COMB.seq:*
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US-09-985-678-211824
US-09-985-678-211824
US-09-9815-264-28205
US-09-9815-264-23340
US-09-815-264-23319
US-09-815-264-23140
US-09-815-264-23140
US-09-815-264-23140
US-09-815-264-23140
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US-09-815-264-23140
US-09-815-264-766882
US-09-815-264-76889
US-09-815-264-76889
US-09-815-264-76833
US-09-815-264-76830
US-09-815-264-76830
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US-09-815-264-76830
US-09-815-264-798808
US-09-815-264-719446
US-09-815-264-719446
US-09-815-264-719466
US-09-985-678-2798809
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                                                                                                   211824, Ap. Be 2737, Ap. Be 27346, Ap. Be 23146, Ap. Be 23149, Ap. Be 23250, Ap. Be 23
109466,
279880,
18469, 1
238, App
2094, Ap
9, A
App
1, Ap
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US-09-620-394B-4668

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FEATURE: NAME/KEY: misc_feature LOCATION: 1..1069 OTHER INFORMATION: any n = a, g, c, t, ur NAME/KEY: misc_feature LOCATION: 1..1069 OTHER INFORMATION: Ceres Seq. ID 1393173 US-09-620-394B-4668 Sequence 4668, Application US/09620394B GENERAL INFORMATION: APPLICANT: ALEXANDROV, Nickola1 APPLICANT: BROVER, Vyacheslav TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptid TITLE OF INVENTION: Thereby FILE REFERENCE: 2750-1067P US-09-924-035A-520 Sequence 520, Application US/09924035A GENERAL INFORMATION: APPLICANT: Grlach, Jrn TITLE OF INVENTION: Expressed Sequences of Arabidopsis TITLE OF INVENTION: thallana FILE REFERENCE: 2011US 밁 δ CURRENT APPLICATION NUMBER: US/09/620,394B CURRENT FILING DATE: 2000-07-21 NUMBER OF SEQ ID NOS: 9131 SEQ ID NO 4668 LENGTH: 1069 Query Match Best Local : Matches CURRENT APPLICATION NUMBER: US/09/924,035A CURRENT FILING DATE: 2000-08-11 PRIOR APPLICATION NUMBER: US 60/148,784 PRIOR FILING DATE: 1999-08-13 NUMBER OF SEQ ID NOS: 900 TYPE: DNA ORGANISM: Arabidopsis thaliana 690 tgttggttcacaagatttgaca Local 1 tgttggatcacaagatttgata l Similarity 20; Conser Conservative 85.5%; 90.9%; 22 0, Score 18.8; I Pred. No. 16; 0; Mismatches t, unknown, DΒ 9 2 or other Length 1069; Indels 0, Gaps 0,

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Sequence 21824, Application US/09985678
GENERAL INFORMATION:
APPLICANT: Cheikh, Nordine
APPLICANT: Cheikh, Nordine
TITLE OF INVENTION: Annotated Plant Genes
FILE REFERENCE: 16517.255/38-21(15097)F
CURRENT APPLICATION NUMBER: US/09/985,678
CURRENT FILING DATE: 2001-11-05
PRIOR APPLICATION NUMBER: US 09/304,517
PRIOR FILING DATE: 1999-05-06
NUMBER OF SEQ ID NOS: 295529
SEQ ID NO 211824
LENGTH: 622
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NAME/KEY: misc_feature

LOCATION: (1)...(249)

OTHER INFORMATION: n = A,T,C

US-09-924-035A-520
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; Sequence 2773, Application US/09922279A
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SEQ ID NO 520
LENGTH: 249
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Best Local Similarity
Matches 19; Conserv
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Best Local Similarity
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SEQ
                                                                            APPLICANT: Drmanac, Radoje T.
TITLE OF INVENTION: Novel Contigs Obtained
TITLE OF INVENTION: From Various Libraries
FILE REFERENCE: 785
CURRENT APPLICATION NUMBER: US/09/922,279A
CURRENT FILING DATE: 2001-08-03
SOFTWARE:
SEQ ID NO 2
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                                              PRIOR FILING DATE: 2000-01-25
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                                 NUMBER OF SEQ ID NOS:
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                                                                                                                                                                                                                                                                                                                                  520 tgatggatctcaagctttgata
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: pt_SP_genes Version 1.0
2773
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Pred. No. 81;
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Pred. No. 8
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US-09-922-279A-2773
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US-09-815-264-82805
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US-09-815-264-82805
                                                                                                                                                                               US-09-982-402-1036
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Matches 19; Conserv
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SEQ ID NO 82805
LENGTH: 46891
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Best Local Similarity
Matches 19; Conserv
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CURRENT FILING DATE: 2001-03-23
PRIOR APPLICATION NUMBER: US 09/620,392
PRIOR FILING DATE: 2000-07-19
PRIOR APPLICATION NUMBER: US 09/702,134
PRIOR FILING DATE: 2000-10-31
PRIOR FILING DATE: 2000-10-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Wu, Wei TITLE OF INVENTION: Genomic Plant Sequences And Uses Thereof FILE REFERENCE: 38-21(51237)G
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                                                 APPLICANT: [list inventors here]
TITLE OF INVENTION: [list title here]
FILE REFERENCE: [list docket ID here]
CURRENT APPLICATION NUMBER: US/09/982,402
CURRENT FILING DATE: 2001-10-17
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ORGANISM: Homo sapiens
FEATURE:
                            NUMBER OF SEQ ID NOS: 3553
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SOFTWARE: PERL Program SEQ ID NO 1036
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Kovalic, David K.
Liu, Jingdong
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on, Stanton B.
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Pred. No. 1.3e
0; Mismatches
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; ORGANISM: Zea mays; FEATURE: 
; NAME/KEY: misc_feature
; OTHER INFORMATION: Incyte ID No: 700196524H1
US-09-982-402-1036
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US-09-815-264-26466/c
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Best Local Similarity
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APPLICANT:
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Best Local Similarity
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APPLICANT: Liu, Jingdong
APPLICANT: McIninch, James
APPLICANT: Wu, Mei
TITLE OF INVENTION: Genomic Plant Sequences and Uses Thereof
FILE REFERENCE: 38-21(51237)G
CURRENT APPLICATION NUMBER: US/09/815,264
CURRENT FILING DATE: 2001-03-23
PRIOR APPLICATION NUMBER: US 09/620,392
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Ruff, Thomas G.
APPLICANT: Shukla, Hridayabhiranjan
TITLE OF INVENTION: Nucleic Acid Molecules And Other Molecules Associated With
TITLE OF INVENTION: Plants
TITLE OF INVENTION: Plants
FILE REFERENCE: 16517.250[38-21(51930)B]
CURRENT EPPLICATION NUMBER: US/09/849,526A
CURRENT FILING DATE: 2001-05-07
                                                                                                                                                                                APPLICANT: Boukharov, Andrey A. APPLICANT: Cao, Yongwei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA
TYPE: DNA
ORGANISM: Zea mays subsp. mexicana
FEATURE:
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Koshi, Jeffrey M.
Kovalic, David K.
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90.08;
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Pred. No. 1.3e
0; Mismatches
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; PRIOR FILING DATE: 2000-07-19
; PRIOR APPLICATION NUMBER: US 09/702,134
; PRIOR FILING DATE: 2000-10-31
; NUMBER OF SEQ ID NOS: 109669
; SEQ ID NO 26466
; EENCTH: 849
; TYPE: DNA
; ORGANISM: Oryza sativa
US-09-815-264-26466
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US-09-815-264-63319
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LENGTH: 1046
TYPE: DNA
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Best Local Similarity 90.0%;
Matches 18; Conservative
                                                                                                                                                                                                                                    Sequence 91672, Appl GENERAL INFORMATION:
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Best Local Similarity
Matches 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 63319, Application US/09815264
APPLICANT: Wu, Wei APPLICANT: Wu, Wei TITLE OF INVENTION: Genomic Plant Sequences FILE REFERENCE: 38-21(51237)G CURRENT APPLICATION NUMBER: US/09/815,264
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APPLICANT: Cao, Yongwei
APPLICANT: Dotson, Stanton B.
APPLICANT: Koshi, Jeffrey M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: US 09/620,392
PRIOR FILING DATE: 2000-07-19
PRIOR APPLICATION NUMBER: US 09/702,134
PRIOR FILING DATE: 2000-10-31
NUMBER OF SEQ ID NOS: 109669
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Wu, Wei
TITLE OF INVENTION: Genomic Plant Sequences And Uses Thereof
FILE REFERENCE: 38-21($1237)G
CURRENT APPLICATION NUMBER: US/09/815,264
CURRENT FILING DATE: 2001-03-23
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Koshi, Jeffrey M.
Kovalic, David K.
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Koshi, Jeffrey M.
Kovalic, David K.
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Dotson, Stanton B.
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                                                                                                                        Liu, Jingdong
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Pred. No. 1.
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US-09-815-264-46882; Sequence 46882; Application US/09815264; GENERAL INFORMATION:
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US-09-815-264-23140
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US-09-815-264-91672
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SEQ ID NO 23140
LENGTH: 1501
TYPE: DNA
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SEQ ID NO 91672
LENGTH: 1246
TYPE: DNA
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CURRENT APPLICATION NUMBER: US/09/815,264
CURRENT FILING DATE: 2001-03-23
PRIOR APPLICATION NUMBER: US 09/620,392
PRIOR FILING DATE: 2000-07-19
PRIOR APPLICATION NUMBER: US 09/702,134
PRIOR APPLICATION NUMBER: US 09/702,134
PRIOR FILING DATE: 2000-10-31
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Best Local Similarity
                                                                                                                        APPLICANT: Boukharov, Andrey A.
APPLICANT: Cao, Yongwei
APPLICANT: Dotson, Stanton B.
CURRENT APPLICATION NUMBER: US/09/815,264
           TITLE OF INVENTION: Genomic Plant Sequences And Uses Thereof FILE REFERENCE: 38-21(51237)G
                                                                                             APPLICANT:
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                                                                                                            Dotson, Stanton B. Koshi, Jeffrey M.
                                                                                             Kovalic, David K.
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                                                           Liu, Jingdong
McIninch, James
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90.08;
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Pred. No. 1
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Pred. No. 1
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US-09-815-264-76889; Sequence 76889; A
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US-09-815-264-53250
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Best Local Similarity
Thes 18; Conserv.
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CURRENT APPLICATION NUMBER: US/99/815,264
CURRENT FILING DATE: 2001-03-23
PRIOR APPLICATION NUMBER: US 09/620,392
PRIOR TILING DATE: 2000-07-19
PRIOR APPLICATION NUMBER: US 09/702,134
PRIOR APPLICATION NUMBER: US 09/702,134
PRIOR APPLICATION NUMBER: US 09/702,134
PRIOR FILING DATE: 2000-10-31
NUMBER OF SEQ ID NOS: 109669
SEQ ID NO 53250
LENGTH: 1501
                                                                                                 Sequence 76889, Appl GENERAL INFORMATION:
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PRIOR FILING DATE: 2000-10-31
NUMBER OF SEQ ID NOS: 109669
SEQ ID NO 46882
LENGTH: 1501
TYPE: DNA
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Best Local Similarity
APPLICANT:
                                                APPLICANT:
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                                                                              APPLICANT: Boukharov, Andrey A.
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PRIOR APPLICATION NUMBER: US 09/620,392
PRIOR FILING DATE: 2000-07-19
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OTHER INFORMATION: unsure at all n locations
                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Oryza sativa
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          Dotson, Stanton B
Koshi, Jeffrey M.
Kovalic, David K.
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Koshi, Jeffrey M.
Kovalic, David K.
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                                                           Cao, Yongwei
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                                                                                                                 , Application US/09815264
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Jingdong
                                               Stanton B.
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90.0%;
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Pred. No. 1.4e+02;
Pred. No. 1.4e+02;
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Pred. No. 1.4e+02;
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APPLICANT: Wo, Wei

APPLICANT: Wo, Wei

TITLE OF INVENTION: Genomic Plant Sequences And Uses Thereof
FILE REFERENCE: 38-21(51237)G

CURRENT APPLICATION NUMBER: US/09/815,264

CURRENT FILING DATE: 2001-03-23

PRIOR APPLICATION NUMBER: US 09/620,392

PRIOR FILING DATE: 2000-07-19

PRIOR FILING DATE: 2000-07-19

PRIOR APPLICATION NUMBER: US 09/702,134

PRIOR APPLICATION NUMBER: US 09/702,134

PRIOR FILING DATE: 2000-10-31

NUMBER OF SEQ ID NOS: 109669

SEQ ID NO 76889

LENGTH: 6940

TYPE: DNA

ORANISM: OTYZA SATIVA

FEATURE:
NAME/KEY: UNSUTE

LOCATION: (1)..(6940)

OTHER INFORMATION: unsure at all n locations

US-09-815-264-76889
                                                                                                                                                                                                                                             APPLICANT: McIninch, James

APPLICANT: Wu, Wei

APPLICANT: Wu Wei

TITLE OF INVENTION: Genomic Plant Sequences And Uses Thereof

FILE REFERENCE: 38-21(51237)G

CURRENT APPLICATION NUMBER: US/09/815,264

CURRENT FILING DATE: 2001-03-23

PRIOR APPLICATION NUMBER: US 09/620,392

PRIOR FILING DATE: 2000-07-19

PRIOR APPLICATION NUMBER: US 09/702,134

PRIOR APPLICATION NUMBER: US 09/702,134

PRIOR FILING DATE: 2000-10-31

NUMBER OF SEQ ID NOS: 109669

SEQ ID NO 81045

LENGTH: 12081

TYPE: DNA

CORANISM: OTYZA Sativa

US-09-815-264-81045
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APPLICANT: Cao, Yongwei
APPLICANT: Dotson, Stanton B.
APPLICANT: Koshi, Jeffrey M.
APPLICANT: Kovalic, David K.
APPLICANT: Liu, Jingdong
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                                                                                                                                                                                           Query Match 76.4%;
Best Local Similarity 90.0%;
Matches 18; Conservative
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3165 ttggaccaaaagatttgata 3184
                                                                                                     Dotson, Stanton B.
Koshi, Jeffrey M.
Kovalic, David K.
Liu, Jingdong
McIninch, James
                                                                                                                                                                                             Score 16.8; DB 6;
Pred. No. 1.7e+02;
0; Mismatches 2;
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Run

U60594 Leptospira Description

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194471 Homo sapi
194471 Homo sapi
194471 Homo sapi
194471 Leptosphil
194471 Leptos

JOURNAL MEDLINE REFERENCE AUTHORS 2 (bases 1 to 1481)
Adler, B., Chappel, R.J., Barranton, G., Bulach, D.M.,
Billinghurst, M.L., Letocart, M., Merlen, F., Serrano, M.S.
Perolat, P. Bacteria; Spirochaetales; Leptospiraceae; Leptospira.

1 (bases 1 to 1481)
Perolat, P., Chappel, R.J., Adler, B., Baranton, G., Bulach, D.M.,
Billinghurst, M.L., Letocart, M., Merien, F. and Serrano, M.S.
Leptospira fainei sp. nov., isolated from pigs in Australia
Int. J. Syst. Bacteriol. 48 Pt 3, 851-858 (1998) gene, partial sequence.

BCT

10-SEP-1998

SUMMARIES

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REFERENCE
AUTHORS
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Leptospira
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Submitted (12-JUN-1996) Microbiology, Monash University,
Rd., Clayton, VIC 3168, Australia
                                                                                                                                                                                                                                                                                                                                                                                                                     Shin, M.-S.
Direct Submission
Submitted (05-SEP-1997) to the DDBJ/EMBL/GenBank databases. Mi-Su
Submitted (05-SEP-1997) to the DDBJ/EMBL/GenBank databases. Mi-Su
Shin, Laboratory of Marine Molecular Microbiology, Faculty of
Shin, Laboratory of Marine Molecular Microbiology, Faculty of
Agriculture, Kyoto University; Olwakecho, Kitashirakawa, Sakyoku,
Kyoto, Kyoto 306-01, Japan (E-mail:misun@kais.kyoto-u.ac.jp,
Tel:075-753-6224, Fax:075-753-6226)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Spirochaeta sp.
Bacteria; Spirochaetaceae;
1 (bases 1 to 354)
                                                                                                                                       l Similarity 95.5
21; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Spirochaeta sp. (s
strain:FO-95) DNA.
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/strain="F0-95"
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<1. >354
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/db_xref="taxon:48782"
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Bacteria; Spirochaetales; I
1 (bases 1 to 283)
Woo,T.H.S., Patel,B.K.C., (
Norris,M. and Dohnt,M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            sequence.
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Location/Qualifiers
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Woo,T.H.S., Patel,B.K.C.,
Norris,M. and Dohnt,M.
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Leptospira inadai
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/strain="48/95"
/db_xref="taxon:173"
                                                                                                                                                   QLD 4111, Australia
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /product="16S rRNA"
60 c 88 g
                                                                                                 /db_xref="taxon:29506"
                                                                                                               /organism="Leptospira
/strain="79/95"
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         85.5%;
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        Score 18
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Bacteria; Spirochaetales; Leptospiraceae; Leptospira.

1 (bases 1 to 288)
Woo,T.H.S., Smythe,L.D., Symonds,M., Norris,M., Dohnt,M.,
Brenner,D.J. and Patel,B.K.C.
Identification of Leptospira inadai by continuously monit
                                                                                                                                                                WOO,T.H.S., Smythe,L.D., Symonds,M., Brenner,D.J. and Patel,B.K.C. Direct Submission
/product="16S ribosomal RNA"
61 c 88 g 62 t
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Pred. No. 78;
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                                                                                                                                                  Science,
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                                                                                                                                                                                                                                                                   continuously monitoring
                                                                                                                                                                                                                                                                                                                                                                                                                                      partial sequence.
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Direct Submission
Submitted (24-MAR-1997) School
Brisbane, QLD 4111, Australia
Location/Qualifiers
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               Direct Submission
Submitted (24-MAR-1997) School of
Brisbane, QLD 4111, Australia
                                                     2 (bases 1 to 288)
Woo,T.H.S., Smythe,L.D., Symonds,M.,
Brenner,D.J. and Patel,B.K.C.
                                                                                                         Woo,T.H.S., Smythe,L.D., Symonds,M., Norris,M., Dohnt,M., Brenner,D.J. and Patel,B.K.C.
Identification of Leptospira inadai by continuously monit fluorescence during rapid cycle PCR
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/strain="68/94"
/db_xref="taxon:29506"
<1. .>288
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Pred. No. 78;
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U94977
Leptospira inadai.
Leptospira inadai
Bacteria; Spirochaetales; Leptospiraceae; Leptospira.
1 (bases 1 to 288)
Moo,T.H.S., Smythe,L.D., Symonds,M., Norris,M., Dohnt
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Woo,T.H.S., Smythe,L.D., Symonds,M.,
Brenner,D.J. and Patel,B.K.C.
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/strain="346/95"
/db_xref="taxon:29506"
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61 c 88 g 62 t
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61 c 88 g 62 t
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DEFINITION ACCESSION

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AUTHORS

JOURNAL REFERENCE

AUTHORS

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U94979
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Brisbane, QLD 4111, Australia
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Brenner, D.J. and Patel, B.K.C. Identification of Leptospira inadai by continuously monitoring fluorescence during rapid cycle PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unpublished
2 (bases 1
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                                                                                     20;
                                                                                                                                                                                                                                                                                                  2 (bases 1 to 288)
Woo,T.H.S., Smythe,L.D., Symonds,M.,
Brenner,D.J. and Patel,B.K.C.
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(bases 1 to 288)
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/strain="1078 VRI"
/db_xref="taxon:29506"
<1...>288
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                                                                                                                                                               /product="16S ribosomal RNA"
61 c 88 g 62 t
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61 c 88 g 62 t
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/strain="218/95"
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QLD 4111, Australia
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Pred. No. 78;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Spirochaeta sp.
Bacteria; Spirochaetales; Spirochaetaceae;
1 (bases 1 to 353)
                                                                                                                                                                                                                               Arabidopsis thaliana cDNA to mRNA.
Arabidopsis thaliana
Arabidopsis thaliana
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
1 (bases 1 to 1012)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Shin,M., Yoshinaga,I., Uchida,A. and Ishida,Y. Phylogenetic analysis by 16S rRNA gene sequencing oligotrophs isolated from the northern basin of La (Mesotrophic Lake)
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                                                                                                                                                   RNS2: a senescence-associated RNase of Arabidopsis that diverged from the S-RNases before speciation Proc. Natl. Acad. Sci. U.S.A. 90 (11), 5118-5122 (1993)
                                                                                                                                                                                                     Green, P.J.
                                                                                                                                                                                                                  Taylor, C.B., Bariola, P.A., delCardayre, S.B.,
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<1. .>353
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80 c 108 g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /organism="Spirochaeta
/strain="SO-104"
                              /gene="RNS2"
16..795
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /sub_species="Freshwater obligate oligotroph"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
                                                                       /organism="Arabidopsis thaliana"
/db_xref="taxon:3702"
                                                                                                                       Location/Qualifiers
/codon_start=1
            /gene="RNS2"
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of Lake Biwa
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20; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                   Submitted (01-MAR-2000) Avila C., Biologia Molecular y Bioquimica, Facultad de Ciencias, E-29071, Malaga, SPAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GDP-mannose pyrophosphorylase; enzyme; vacuolar ribonuclease.
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AJ275979.1 GI:13509286
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Direct Submission
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                                                                                                                                                                     /product="GDP-mannose pyrophosphorylase"
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GVDEVVLAINYQPEVMLNFLKOPETKLEIKITCSQETEPLGTAGPLALARDKLLDGSG
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                                             /number=1
1597. .16
                                                                                                                                          DIGQPRDYITGLRLYLDSLRKKSPAKLTSGPHIVGNVLVDETATIGEGCLIGPDVAIG
PGCIVESGVRLSRCTVMRGVRIKKHACISSSIIGWHSTVGQWARIENMTILGEDVHVS
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/protein_id="AAA51406.1"
/db_xref="GI:289210"
             1660.
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                                           Direct Submission
Submitted (09-MAR-2000) The Institute for Genomic Research, 9712
Medical Center Dr., Rockville, MD 20850, USA
On Dec 17, 1999 this sequence version replaced gi:2642152.
The sequence and annotation of chromosome 2 were merged from those of the individual clones on this chromosome after removing
                                                                                                                                                                                                                                                                                                                                                                              Arabidopsis thaliana
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.

1 (bases 1 to 72590)
Lin,X., Kaul,S., Rounsley,S.D., Shea,T.P., Benito,M.-I., Town,C.D.,
Fujii,C.Y., Mason,T.M., Bowman,C.L., Barnstead,M.E.,
Fujii,C.Y., Mason,T.M., Bowman,C.L., Barnstead,M.E.,
Koo,H., Moffat,K.S., Cronin,L.A., Shen,M., Vanaken,S.E., Umayam,L.,
Koo,H., Moffat,K.S., Cronin,L.A., Shen,M., Vanaken,S.E., Umayam,L.,
Tallon,L.J., Gill,J.E., Adams,M.D., Carrera,A.J., Creasy,T.H.,
Goodman,H.M., Somerville,C.R., Copenhaver,G.P., Preuss,D.,
Nlerman,W.C., White,O., Eisen,J.A., Salzberg,S.L., Fraser,C.M. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     complete sequence. Sequence from clones T517 AC003000 AE002093 AC003000.2 GI:6598383
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Arabidopsis thaliana chromosome II section
                                                                                                                                                                                                                                                                     Nature 402 (6763),
20083487
                                                                                                                                                                                                                                                                                                                                          Sequence and analysis of chromosome
                                                                                                                                                                                                                                                                                                                                                                      Venter, J.C.
                      overlaps. For detailed information, please see the TIGR web site
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(http://www.tigr.org/tdb/at/at.html).
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/protein_id="CAC35357.1"
/db_xref="gi:13599289"
/translation="MRGKGQVVDGDNGIVYTHQDIPIQA"
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/protein_id="CAC35356.1"
/db_xref="GI:13509288"
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3288. .3473
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        We thank the CSHL/WashU/ABI consortium for sequencing BAC clones F6P23, F5J6, T17A5, and T13L16, the ESSA group for sequencing clone F13D4, and Scott Jackson, Jiming Jiang, Klaus Meyer, Eric Richards and Satoshi Tabata for helpful assistance. In addition, we would like to thank the TIGR Bioinformatics Department, especially Lixin Zhou, Hanif Khalak, Michael E. Heaney, Lily Fu, Feng Liang, Jeremy Peterson, Michael Holmes, and Delwood Richardson for software and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              http://genome.wustl.edu/edgy/tRNAscaar-SE/). Simple repeats were identified by repeatmasker (Arian Smit, http://ftp.genome.washington.edu/RM/RepeatMasker.html). Genes are numbered from the top; to bottom of the chromosomo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (ftp://arthur.epm.ornl.gov/pub/xgrail), Genefinder (Phil Gre
University of Washington), Genscan (Chris Burge,
http://gnomic.stanford.edu/GENSCANW.html), and NetPlantGene
(http://www.cbs.dtu.dk/services/NetGene2/), searches of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           prediction programs including GRAIL (ftp://arthur.epm.ornl.gov/pub/xgrail), Genefinder (Phil Green,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Genes were identified by a combination of three methods: Gene
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/rpt_family="POLY_A"

complement(<4584...>5909)

/gene="At7339710"

/gene="At7339710"

/note="T517.1; similar to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /#UDIC="T517.1; Similar to GP|2245012|gn1|PID|e327000|297341" complement(4584, .5909) /gene="At2g39710" /note=""."
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/rpt_family="POLY_A"
complement(1266..1291)
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/POOLEIn_id="AABB7120
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                                                                                                                             DLCYKVGSTTRPNFSGLPMVSLMFRGAEMSVSGQKLLYRVNGAGSEGKEEVYCFTFGN
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/chromosome="II"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /organism="Arabidopsis thaliana"
/cultivar="Columbia"
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/codon_start=1
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/gene="At2g39720"
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GTHHAVLSSYEYVSQGLRQYNLDNMMDGFYIAPAFMDKLVVHIFKNFLFLRNIKVPLIK
LGIMGGKGGKSFQCELVMAKMGINPIMMSAGELESGNAGEPAKLIRQRYREAADLIK
KGKMCCLFINDLDAGAGRAGGTTGYTVNNQMVNATLMNIADNPTNVQLPGMYNKEENA
RVPICTGNDFSTLYAPLIROGRMEKEYWAPTREDRIOVKGFIRTDKIKDEDIVTLV
DQFPGGSIDFFGALRARVYDDEVRKEVESLGVEKIGRILVNSREGPFWFEGPEMTYEK
LORFGGNHLVMEGENVKRVQLAETYLSQAALGDANADAIGRGTFYGKGAQQVNLPVPEG
CTDPVARNEDPTARSDDGTCVYNF"
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complement()oin(13323. .13599,13724. .14005,14087. .14559,
14651. .14736,14836. .14941,15033. .15349,15834. .16055))
/gene="At2339730"
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SETTITTOHRSPTRFPPDSSSSSTPSASMHADNSPTPTIVTRTRSNRSPNPVIVLRG
SAAAPSSDVYSEGLDRSAFGMYYDDGTDSGLBFLPPSMTEFLLGSGFDRLLDQISQIE
LNTNRNLRSCEHPPASKSAIEALPLIEIDPTHLLSDSQSHCAVCKENFVLKSSAREMP
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14651...14736,14836...14941,15033...15349,15834...1
/gene-"At2g39730"
complement(13323...16055)
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IWRLPGGGFAVGRIPGGWRGGDRMMPVYYTEVDGGRLGDERLPRRVAWGSRRGGRDGG
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complement(join(13484. .13599,13724. .14005,14087. .14559,
14651. .14736,14836. .14941,15033. .15349,15834. .15878))
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/db_xref="GI:2642154"
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/gene="At2g39720"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AQSINKKSINGSEKVLAVKEDKQTDGDRWRGLAYDTSDDQQDITRGKGMVDSVFQAPMGT
GTHHAVLSSVEYVSQGLRQYNLDNMMDGFYIAPAFMDKLVVHITKVRELTLDRIKVPLI
LGIWGGKGQGKSFQCELVMAKMGINPIMMSAGELESGNAGEPAKLIRQRYREAADLIK
KGKMCCLFINDLDAGAGRMGGTTGYTVNNQMVNATLMNIADNPTMYQLDEGMYNKEENA
RVPIICTGNDFSTLYAPLIRDGRMEKFYWAPTREDRIGVCKGIFRTDKIKDEDIVTLV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       complement(join(13579...13599,13713...14005,14087.
14651...14736,14836...14941,15033...15349,15834...
/gene="At2g39730"
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/protein_id="AAB87122.1"
/db_xref="GI:2642155"
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/codon_start=1
             <18783. .>21237
/gene="At2939740"
/note="T517.4; predicted by genscan"
/note="T517.4; predicted by 19352,19496.
join(18783. .18921,19188. .19352,19496.
19965. .20034,20131. .20261,20593. .212
                                                                                                                                                                                                             join(<18783. 18921,19188. .19352,19496. .19674,19766. .19861,
19965. .20034,20131. .20261,20593. .>21237)
/gene="Mc2g39740"
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/note="alternative splice form 1 GP|166834|M86720"
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/note="T517.3"
                                                                                                                                                                                                                                                                                                                                                                                                                                   LMEYGNMLVMEQENVKRVQLAETYLSQAALGDANADAIGRGTFYGKTEEKEPSK"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DQFPGQSIDFFGALRARVYDDEVRKFVESLGVEKIGKRLVNSREGPPVFEQPEMTYEK
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/protein_id="AAB87137.1"
/db_xref="GI:2642170"
/translation="MAAAVSTYGAINRAPLSINGSGSGAVSAPASTFIGKKVVTVSRF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        translation="MAAAVSTVGAINRAPLSLNGSGSGAVSAPASTFLGKKVVTVSRF/
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                                                                                                                                                                                                                                                                                                           /rpt_family="(TA)n"
join(<18783
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note="alternative splice form 2 GP|166835|M86720"
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             496. .19674,19766.
.21237)
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AF165146
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Submitted (06-JUL-1999) Genome Analysis, Institute of Molecular Biotechnology, Beutenbergstrasse 11, Jena 07745, Germany On Jun 7, 2001 this sequence version replaced gi:8151945.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 (bases 1 to 121524)
Schilhabel,M.B., Baumgart,C., Blechschmidt,K., Dette,M., Jahn,N.,
Lehmann,R., Menzel,U., Polley,A., Reichwald,K., Schudy,A.,
Siddiqui,R., Taudien,S., Wen,G., Schlegelberger,B., Siebert,R.,
Rosenthal,A. and Platzer,M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AF165146 121524 bp DNA HTG 07-JUN HOMO Sapiens chromosome 8 clone CTA-397H3 map 8q12-8q13, SEQUENCING IN PROGRESS ***, 7 unordered pieces.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Schudy,A., Blechschmidt,K., Schillhabel,M., Baumgart,C., Menzel,U., Weber,J., Schattevoy,R. and Rosenthal,A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Chromosome 8 genomic sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
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NOTE: This is a 'working draft' sequence. It currently
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (bases 1 to 121524)
                                         Center project name: H171
Center clone name: CTA-397H3
Center clone name: CTA-397H3
Center clone name: CTA-397H3
Sequencing vector: M13; 100% of reads
Chemistry: Dye-terminator Big Dye; 100% of reads
Assembly program: Phrap; version 0.990329
Consensus quality: 115981 bases at least Q40
Consensus quality: 117981 bases at least Q30
Consensus quality: 117981 bases at least Q20
Quality coverage: 9.57 in Q20 bases; sum-of-contigs
                                                                                                                                                                                                                                                                                                                Contact: gscj-submit@genome.imb-jena.de
                                                                                                                                                                                                                                                                                                                                                                                                 Center: Insitute of Molecular Biotechnoloy Center code: IMB
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                                                                                                                                                                                                                                                                                                                                                                        Web site: http://genome.imb-jena.de/
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22328. .22585,22834. .22993,23212. .23322,23481.
23847. .23996,24089. .>24814))
/gen="At2g39750"
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/db_xref="GI:2642156"
/db_xref="GI:2642156"
/db_xref="GI:2642156"
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Lrgatyopegsfysnlftrwgdldisvdlfsgssilftgkkokotllghllralrasg
Lwyklofviharvpilkvvsghoriscoltsidnubgllksgrlerbisbldgrfrodyl
Lvkermakahundskygffenystsllvifhpotycyralipplryipyrkavddlfdy
Rktabesiaqvtaaniarfkseraksvnrsslsellysffakvedpfeqpynaarsys
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!-8q13, ***
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AC083961
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                                                                                                                                               Birren, B., Linton, L., Nusbaum, C., Lander, E., Abraham, H., Allen, N., Anderson, S., Barna, N., Bastien, V., Beda, F., Boguslavkiy, L., Boukhgalter, B., Brown, A., Burkett, G., Campopiano, A., Castle, A., Choepel, Y., Colangelo, M., Collins, S., Collymore, A., Cooke, P., DeArellano, K., Dewar, K., Diaz, J.S., Dodge, S., Ferreira, P., DeArellano, K., Gage, D., Galagan, J., Gardyna, S., Ginde, S., Goyette, M., Graham, L., Grand-Pierre, N., Hagos, B., Heaford, A., Horton, L., Graham, L., Grand-Pierre, N., Hagos, B., Heaford, A., Horton, C., Liu, G., Liev, I., Johnson, R., Jones, C., Kann, L., Karatas, A., LaRocque, K., Lamazares, R., Landers, T., Lehoczky, J., Levine, R., Lieu, C., Liu, G., Macdonald, P., Marquis, N., McCarthy, M., McBwan, P., McKernan, K., McPheeters, R., Meldrim, J., Meneus, L., Mihova, T., Mlenga, V., Morrow, T., Mendrim, J., Norman, C.H., O'Connor, T., Morphy, T., Navlor, J., Norman, C.H., O'Connor, T., Miroya, T., Mandon, C.H., O'Connor, T., Morphy, T., Navlor, J., Norman, C.H., O'Connor, T., Miroya, T., Mandon, C.H., O'Connor, T., Miroya, Mandon, Mando
Morrow,J., Murphy,T., Naylor,J., Norman,C.H., O'Connor,T., O'Donnell,P., O'Neil,D., Olivar,T.M., Oliver,J., Peterson,K., Pierre,N., Pisani,C., Pollara,V., Raymond,C., Rieback,M., Riley,R., Rogov,P., Rothman,D., Roy,A., Santos,R., Schauer,S., Severy,P., Sougnez,C., Spencer,B., Stange-Thomann,N., Stojanovic,N.,
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7: gap of unknown length
0: contig of 2243 bp in length
0: gap of unknown length
4: contig of 17994 bp in length
4: gap of unknown length
6: contig of 21932 bp in length
6: gap of unknown length
6: gap of unknown length
13: contig of 17627 bp in length
3: contig of 17627 bp in length
3: gap of unknown length
8: contig of 23835 bp in length
8: gap of unknown length
4: contig of 36106 bp in length.
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Pred. No. 71;
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Strauss, N., Subramanian, A., Talamas, J., Tesfaye, S., Theodore Tirrell, A., Travers, M., Trigilio, J., Vassiliev, H., Viel, R., Wilson, B., Wu, X., Wyman, D., Ye, W.J., Young, G., Zainoun, J., Zimmer, A. and Zody, M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Submitted (08-OCT-2000) Whitehead Institute/MIT Center for Research, 320 Charles Street, Cambridge, MA 02141, USA On Feb 15, 2001 this sequence version replaced gi:10717228.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             All repeats were identified using RepeatMasker:
Smit, A.F.A. & Green, P. (1996-1997)
http://ftp.genome.washington.edu/RM/RepeatMasker.html
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                      20537: gap of 100 bp
21159: contig of 622 bp in length
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3251: gap of 100 bp
9962: contig of 711 bp in length
10062: gap of 100 bp
10776: contig of 714 bp in length
10876: gap of 100 bp
11474: contig of 598 bp in length
11574: gap of 100 bp
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18: gap of

8145: con
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                                                                                                                         739: gap of 100 bp
13952: contig of 1213 bp
052: gap of 100 bp
14838: contig of 786 bp
938: gap of 100 bp
15849: contig of 911 bp
15849: contig of 910 bp
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46344 46443; gap of 100 bp
46444 47374; contig of 1131 bp in le
47575 476774; gap of 100 bp
47675 50217; contig of 2543 bp in le
50218 50317; gap of 100 bp
50318 51338; contig of 1021 bp in le
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2 80685: contig of 3544 b

36 80785: gap of 100 bp

36 80785: gap of 3040 bp

36 83925: gap of 100 bp

36 83925: gap of 100 bp

3721: contig of 3796 b
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62360: gap of 100 bp
64120: contig of 1760 bp
6220: gap of 100 bp
65979: contig of 1759 bp
5079: gap of
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                     92828:
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100 bp
4700: gap of 1022 k
24891: rrr
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37036: contig of 1673 bp in
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138963: contig of 1827 bp in
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23478: contig of
828: gap of
96652: cont
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40807: contig of 1744 bp in
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44598: contig of 1602 bp in
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FEATURES
source
Search completed: December 15, 2001, 03:20:02
Job time: 7692 sec
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199907 100006: gap of 100 bp
100007 102910: contig of 2904 bp in length
102911 103010: gap of 100 bp
103011 107188: contig of 4178 bp in length
107189 107288: gap of 100 bp
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111642 121990: contig of 5149 bp in length
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126622: contig of 4532 bp in length
126623 126792: gap of 100 bp
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126723 129090: contig of 2368 bp in length
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/organism="Homo sapiens"
//bb_xref="taxon:9606"
//clone="RP11-22E14"
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C. elegans inhibit	AAT72715	18	1075	0	15.6	45	()
Adenovirus SCAR.RG	AAV72026	20	834	0	٠	44	4.3
Adenovirus pACTSG2	AAV72025	20	795	0	٠	43	(3
Human olfactory re	AAH31988	22	635	0.	15.6	42	O
S. epidermidis ope	AAH53651	22	452	0	•	41	
Human olfactory re	AAH32175	22	342	0	15.6	40	(3
EST clone HK650.	AAV88753	20	297	0	15.6	39	()
Human hypoxia resp	AAZ51556	21	4146	۳.	15.8	38	
H. pylori GHPO 131	AAX14095	19	961	۲.	15.8	37	
Human gene express	AAZ15467	20	843	۳.	15.8	36	
Arabidopsis thalia	AAC51908	21	740	71.8	15.8	35	1,1
Human secreted pro	AAC10808	21	736	۲.	15.8	34	
Human secreted pro	AAC09187	21	670	۳.	15.8	33	1,
	AAH32060	22	467	۲.	15.8	32	u
Potato pollen cell	AAV59091	19	10811	ω	16.2	31	12
Maize acetyl CoA c	AAZ49820	21	7470	ω.	16.2	30	
	AAV29317	19	7470	Ψ	16.2	29	
Maize acetyl CoA c	AAT39905	17	7470	ω.	16.2	28	
Potato subclone pG	AAA39732	21	5400	ω.	16.2	27	12
A3 maize ACCase cD	AAQ42933	14	4346	ω.	16.2	26	
M. leprae gyrA pre	AAT64683	18	3822	Ψ	16.2	25	
EcoRI fragment of	AAZ49816	21	2001	ω.	16.2	24	
Maize acetyl CoA c	AAT39904	17	2001	73.6	16.2	23	
	AAT17111	17	2000	ω ·	16.2	22	
(D	AAX61757	20	1473	w	16.2	21	•
Arabidopsis thalia	AAC39657	21	1452	ω.	16.2	20	•
	AAC49931	21	1450	ω.	16.2	19	13
	AAH33007	22	1376	ω.	16.2	18	•
	AAV59636	19	1291	ω.	16.2	17	
	AAC59054	21	1250	ω.	16.2	16	()
Human secreted pro	AAV84598	20	951	ω.	16.2	15	()
-	AAH33702	22	335	ω ·	16.2	14	()
yptus	AAC56059	21	2444	٠.	16.4	13	
Maize male fertili	AAT90543	18	2582	76.4	16.8	12	• 2

ALIGNMENTS

(first entry)

22 ВP

New isolated pathogenic Leptospira bacterium - useful for, edeveloping products for conferring protective immunity, and forophylactic or therapeutic treatment Chappel RJ 07-MAR-1997; Infection; pathogenic Leptospira; protective immunity; therapy;
diagnosis; ss. WPI; 1998-520791/44 (AGRI-) AGRIC VICTORIA SERVICES PTY LTD. (PIGR-) PIG RES & DEV CORP. 06-MAR-1998; 17-SEP-1998. WO9840099-A1. Leptospira sp. 98WO-AU00145. 97AU-0005494. for e.e

8 x 5 5 5 5 5 5 5 5 5 5 8 x 8

Sequence 22 BP; 7 A; 2 C;

5 G; 8 T; 0 other;

.08

DB 19; 0.11;

Length

22;

Claim

15;

Page 72;

94pp; English

This sequence represents a Leptospira DNA sequence isolated from the pathogenic Leptospira (LS) bacterium of the invention. The bacterium belongs to serogroup Hurstbridge or serovar hurstbridge or the species L. fainei. The LS bacteria can becused for conferring protective immunity against pathogenic LS bacteria in humans or animals. The bacteria can also be used for prophylactic or therapeutic treatment of infections. The DNAs and antibodies may also be used for detection and diagnosis of past or present LS infection.

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Best Local S
Matches 22
Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                           AAV58897;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAV58897 standard;
                                                                This sequence represents a Leptospira DNA sequence isolated from the pathogenic Leptospira (LS) bacterium of the invention. The bacterium belongs to serogroup Hurstbridge or serovar hurstbridge or the species L. fainei. The LS bacteria can be used for conferring protective immunity against pathogenic LS bacteria in humans or animals. The bacteria can also be used for prophylactic or therapeutic treatment of infections. The DNAs and antibodies may also be used for detection and
                                                                                                                                                                New isolated pathogenic Leptospira bacterium - useful for developing products for conferring protective immunity, prophylactic or therapeutic treatment
                                                                                                                                                                                                                          Chappel
                                                                                                                                                                                                                                                                                              06-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                    Leptospira fainei.
                                     Sequence 22
                                                                                                                                               Claim 15;
                                                                                                                                                                                                        WPI; 1998-520791/44.
                                                                                                                                                                                                                                             (PIGR-)
                                                                                                                                                                                                                                                       (AGRI-)
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                                                                                                                                               Page 70; 94pp; English.
                                                                                                                                                                                                                                                                                                                                                                         pathogenic Leptospira; protective immunity; therapy;
ss.
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                                       BP; 7 A; 2 C; 5
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diagnosis;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This sequence represents a Leptospira DNA sequence isolated from the pathogenic Leptospira (LS) bacterium of the invention. The bacterium belongs to serogroup Hurstbridge or serovar hurstbridge or the species L. fainei. The LS bacteria can be used for conferring protective immunity against pathogenic LS bacteria in humans or animals. The bacteria can also be used for prophylactic or therapeutic treatment of infections. The DNAs and antibodies may also be used for detection and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New isolated pathogenic Leptospira bacterium - useful for, developing products for conferring protective immunity, and prophylactic or therapeutic treatment
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 1477 BP; 390 A; 334 C; 439 G; 314 T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 15;
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  Plant PrAG1
                                                22-JAN-2001
                                                                                                                                           AAA99365 standard;
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promoter related
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                                                                                                                                                                                                                                                                                                                                                              This invention relates to a novel plant promoter gene. The promoter is located in plant reproductive tissue, and the invention includes transgenic plants containing the promoter. The promoter can be used to produce plants which have a diminished reproductive capacity or which are sterile. The constructs can also be used to transform agronomically important plants in which modulation of reproductive capacity (particularly the timing and abundance of flowering) is desirable, e.g. cereals, rice, maize, wheat, barley, oats, rye, soybean and canola. The present sequence represents DNA encoding the plant reproductive promoter (PrAGI) of the invention.
                                                                                                 Hybridisation assay; genetic mapping; gene expression control; protein identification; signal transduction pathway; metabolic pathway; promoter; termination sequence; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 15; Page 42-43;
                    25-FEB-2000;
                                         06-SEP-2000.
                                                            EP1033405-A2.
                                                                                Arabidopsis thaliana.
                                                                                                                                          Arabidopsis thaliana DNA fragment SEQ
                                                                                                                                                              17-OCT-2000
                                                                                                                                                                                                    AAC37325 standard; DNA; 1069
                                                                                                                                                                                                                                                                                                                                            Sequence 1012 BP; 287 A; 197 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel plant reproductive tissue promoter, useful to produce plants which have a diminished reproductive capacity or which are sterile
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (CART-) CARTER HOLT HARVEY LTD.
(TASM-) TASMAN BIOTECHNOLOGY LTD.
(UNMT) UNIV MICHIGAN TECHNOLOGICAL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17-MAR-1999;
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  25-FEB-1999;
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Similarity 90.9%;
20; Conservative
                      2000EP-0301439
                                                                                                                                                             (first entry)
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  99US-0121825
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11-MAY-1999
11-JUN-1999
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9908-0125264
9908-0126765
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RESULT 6
AAA79696/c
ID AAA79696;
XX AAA79696;
AC AAA79696;
XX Eucalyptus
XX Eucalyptus
XX Eucalyptus
XX Eucalyptus
XX Eucalyptus
XX Eucalyptus
XX Plant cell
KW environmet
KW elongatior
XX Eucalyptus
XX WO2000421:
XX WO2000421:
XX WO2000421:
XX Il-JAN-200
XX Il-JAN-200
XX Il-JAN-199
PR 11-JAN-199
PR 01-NOV-199
PA (GENE-) GI
XX
PI Strabala:
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24-0CT-1999
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28-0CT-1999
28-0CT-1999
                                                                                                                                                                Eucalyptus grandis; Pinus radiata; plant cell signalling; modulation; environmental change; development; elongation; survival; disease resi:
                                                 12-JAN-1999;
01-NOV-1999;
                                                                                                                                                                                                                    Eucalyptus
                                                                                                                                                                                                                                        27-NOV-2000
                                                                                                                                           Eucalyptus grandis.
                                                                                11-JAN-2000;
                                                                                                   20-JUL-2000
                                                                                                                         WO200042171-A1
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                             (GENE-) GENESIS RES & DEV CORP LTD.
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20; Conserv
                                                                                                                                                                                                                                                                                standard; cDNA;
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nilarity 90.9%;
Conservative
                                                                                                                                                                                                                    grandis
                                                                                                                                                                                                                                        (first entry)
                                                                                 2000WO-US00724.
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990S-0158029

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990S-0160741

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990S-0161406

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990S-0161359

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990S-0161932

990S-0161932

990S-0161933
          Nieuwenhuizen NJ;
                                                  99US-0228986.
99US-0162866.
                                                                                                                                                                                                                    cell signalling
                                                                                                                                                                  disease resistance;
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                                                                                                                                                                                                                                                                                                                                                                          pred. No. 5.8; O; Mismatches
                                                                                                                                                                                                                                                                                                                                    711
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                                                                                                                                                                                                                                                                                  ΒP
                                                                                                                                                                  Monterey pine; plant; modification;
transgenic plant; pathogen; growth;
cell proliferation; differentiation;
stance; nutrient metabolism; ss.
                                                                                                                                                                                                                     involved
                                                                                                                                                                                                                                                                                                                                                                                              DB
                                                                                                                                                                                                                     polynucleotide SEQ
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                                                                                                                                                                                                modification;
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14-JUL-1999;
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CC and protein sequences isolated from eucalyptus (Eucalyptus grandis) or
pline (Pinus radiata also known as Monterey pine). The protein sequences
CC are involved in cell signalling. The polynucleotide and protein
CC sequences can be used to modify the response of plant cells to external
CC signals e.g. environmental changes or pathogens during the growth and
CC differentiation, elongation and survival, resistance to disease and
CC differentiation, elongation and survival, resistance to disease and
CC utrient metabolism. Examples of modifications which can be produced are
CC altered fruit ripening and senescence of leaves and flowers e.g. to
CC delay senescence and prolong the life of cut flowers or enhance
CC senescence of reproductive organs to engineer sterile plants. Other
CC corgans providing fruit and vegetables which have a longer shelf life
CC between harvest and consumption, or to decrease branching frequency in
CC forest tree species giving long stretches of valuable knot-free clear
CC wood which can be used in solid timber furniture and veneers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local S
Matches 19
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           Claim 1; SEQ ID 22189;
                                        diagnostic, forensic, gene therapy and
                                                        obtaining
                                                         New nucleic acid that is a 5' expressed sequence tag (5' EST) for obtaining cDNAs and genomic DNAs that correspond to 5'ESTs and fo
                                                                                                                                       Dumas Milne Edwards J,
                                                                                                                                                                                                                                       21-FEB-2000; 2000EP-0200610
                                                                                                                                                                                                                                                                                                       EP1033401-A2
                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                         Human; 5' EST;
                                                                                                                                                                                                                                                                                                                                                                                                                       Human secreted protein 5' EST, SEQ ID NO: 22189.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAC18114 standard; cDNA; 447 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 2084 BP; 502 A; 434 C; 566 G; 582 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 1; Page 408-409;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Isolated polynucleotide encoding a polypeptide involved in cell signaling used for generating transgenic plants with modified responses to external signals -
                                                                                                                                                                        (GEST ) GENSET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 tgttggatcacaagatttgat 21
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                                                                                                                                                                                                                                                                                                                                                                    therapy;
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19; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                   ; expressed sequence tag; secreted protein; {\tt cDNA} isolation; chromosome mapping; ss.
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90.5%;
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        71pp + CD-ROM; English
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Pred. No. 19;
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                                                                                                                                         Giordano
                                          chromosome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DΒ
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                                        mapping procedures
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RESULT 8
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Best Local :
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                      28-APR-1999;
30-APR-1999;
30-APR-1999;
04-MAY-1999;
05-MAY-1999;
06-MAY-1999;
07-MAY-1999;
                                                                                                                                                  06-APR-1999;
08-APR-1999;
16-APR-1999;
19-APR-1999;
21-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present sequence is one of a large number of 5' ESTs derived from mRNAs encoding secreted proteins. No ORF has yet been conclusively identified within the present sequence. The 5' ESTs were prepared from total human RNAs or polyA+ RNAs derived from 30 different tissues. EST sequences usually correspond mainly to the 3' untranslated region (UTR) of the mRNA because they are often obtained from oligo-dT primed cDNA libraries. Such ESTs are not well suited for isolating cDNA sequences derived from the 5' ends of mRNAs and even in those cases where longer cDNA sequences have been obtained, the full 5' UTR is rarely included. 5' ESTs are derived from mRNAs with intact 5' ends and can therefore be used to obtain full length cDNAs and genomic DNAs. 5' ESTs are also used in diagnostic, forensic, gene therapy and chromosome mapping procedures. They are used to obtain upstream regulatory sequences and to design
                                                                                                                                                                                                                 25-MAR-1999;
29-MAR-1999;
01-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                         Hybridisation assay; genetic mapping; gene expression control; protein identification; signal transduction pathway;
                                                                                                                          23-APR-1999;
23-APR-1999;
                                                                                                                                                                                                                                                        23-MAR-1999
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09-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                   Arabidopsis thaliana
                                                                                                                                                                                                                                                                                                                                                                                                                              metabolic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   expression and secretion vectors.
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                                                                                                                                                                                                                                                                                                                         25-FEB-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                           pathway; promoter;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                          2000EP-0301439
         990S-0130077

990S-0130449

990S-0130510

990S-0131449

990S-0132407

990S-0132486

990S-0132486

990S-0132486

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990S-0132485
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99US-0128714.
99US-0129845.
                                                                                                                                                                                                                 99US-0126264.
99US-0126785.
99US-0127462.
                                                                                                                                                                                                                                                                    99US-0123180
99US-0123548
                                                                                                                                                                                                                                                        99US-0125788
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA; 1290
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       78.2%;
86.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22
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                                                                                                                                                                                                                                                                                                                                                                                                                               termination
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Pred. No. 32;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NO:
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RESULT
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AAC46258
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         25-FEB-2000;
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42 TGTTGTAACACAAGATTTGTTA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      thaliana DNA fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first
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9905-0161405.
9905-0161406.
9905-0161359.
9905-0161361.
9905-0161361.
9905-0161920.
9905-0161993.
9905-0161993.
990S-0121825

990S-012548

990S-0125788

990S-0126785

990S-01267462

990S-01267462

990S-01267462

990S-012845

990S-012845

990S-0130047

990S-013049

990S-0131449

990S-0132486

990S-013421

990S-013421

990S-013421
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Pred. No. 36;
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02-AUG-1999; 03-AUG-1999;

DB

Length 1413; Indels

0;

Gaps

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10-AUG-1999
11-AUG-1999
12-AUG-1999
13-AUG-1999
13-AUG-1999
16-AUG-1999
17-AUG-1999
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28-EEP-1999
28-EEP-1999
29-SEP-1999
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21-SEP
                                                                                                                                                                      14-0CT-1999
18-0CT-1999
21-0CT-1999
                                                          22-OCT-1999;
25-OCT-1999;
25-OCT-1999;
25-OCT-1999;
26-OCT-1999;
990S-0147302

990S-0147192

990S-0147260

990S-0147493

990S-0147493

990S-0148341

990S-0148341

990S-0148341

990S-0149368

990S-0149368

990S-0149930

990S-0149930

990S-0149930

990S-01510866

990S-01510866

990S-01510866

990S-01510866

990S-0151080

990S-0151080

990S-0151080

990S-0151930

990S-0152363

990S-0152363

990S-0153758

990S-0153758

990S-0154018

990S-0155659

990S-0155659

990S-0155659

990S-0155713

990S-0155775

990S-0155823

990S-015933

990S-0160814

990S-0161405

990S-0161405

990S-0161405

990S-0161360
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AAS22799
                  В
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PR
       The invention relates to polynucleotides encoding novel human CC proteins or their active domains. The polypeptides, polynucleotides and antibodies raised against the polypeptides are used in a method of CC treatment of a mammal and prevention of disorders caused by the aberrant CC enclecular weight markers, food supplements, and in antibody production. CC molecular weight markers, food supplements, and in antibody production. CC polypeptides are used to identify compounds which bind to the CC production of recombinant proteins, and in gene mapping, in the CC production of recombinant proteins, and in generating anti-seense DNA or CC RNA and in gene therapy. Polypeptides of the invention can be used to target drugs to a tumour, in assays to determine biological activity, to craise antibodies/elicit an immune response, to determine quantitative CC polypeptides of the invention may also be useful in treating platelet colypeptides of the invention may also be useful in treating platelet colypeptides of the invention may also be useful in treating burns, promoting the poliferation, differentiation and survival of stem cells, as a contraceptive, treating osteoporosis and osteoarthritis, anaemia, cc contraceptive, parkinson's and Huntington's diseases, amylotrophic lateral
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28-OCT-1999;
28-OCT-1999;
29-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; novel protein; ss; Antianaemic; osteopathic; antiinflammatory; immunomodulatory; cytostatic; neuroprotective; vulnerary; nootropic; anticonvulsant; antiarthritic; cerebroprotective; antifungal; antivital; antibacterial; antiallergic; dermatological; haemostatic; antiasthmatic; thrombolytic; immunogen; antibody; gene therapy; neurological disorder; parkinson's disease; inflammatory disorder; cancer; asthma; osteoporosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAS22799;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAS22799
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Parkinson's disease; inflammatory distissue regeneration; immune disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tang YT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25-JAN-2001; 2001WO-US02623
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            02-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200155437-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               576 TGTTGTAACACAAGATTTGTTA 555
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Isolated polypeptides useful for treating anti-inflammatory diseases, nervous system disorders, and for regenerating bone and cartilage \dot{\cdot}
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (HYSE-) HYSEQ INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       25-JAN-2000; 2000US-0491404.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; Page 733-734; 894pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    P-PSDB; AAU14494
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2001-451939/48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 tgttggatcacaagatttgata
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cDNA encoding a novel human protein #365.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Liu C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          99US-0161992.
99US-0161993.
99US-0162142.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Drmanac
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             78.2%;
86.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2038
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          꼅
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 17.2;
Pred. No. 36;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
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sclerosis,

stroke,

viral

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RESULT 11
AAH15352/c
ID AAH15352 standard; cDNA;
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                 The present invention describes primer sets for synthesising 5602 full-length cDNAs defined in the specification. Where a primer set comprises: (a) an oligo-dT primer and an oligonucleotide complementary to the complementary strand of a polynucleotide which comprises one of the 5602 nucleotide sequences defined in the specification, where the oligonucleotide comprises at least 15 nucleotides; or (b) a combination of an oligonucleotide comprising a sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                           29-JUL-1999;
27-AUG-1999;
11-JAN-2000;
02-MAY-2000;
             of an oligonucleotide comprising a sequence complementary to the complementary strand of a polynucleotide which comprises a 5' end sequence and an oligonucleotide comprising a sequence complementary to a polynucleotide which comprises a 3'-end sequence, where the oligonucleotide comprises at least 15 nucleotides and the combination of the 5'-end sequence; a selected from those defined in the specification. The primer sets can be used in antisense therapy and in gene therapy. The primers are useful for synthesising polynucleotides,
                                                                                                                                                                                                                                                                                                                                          WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1929
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   fungal infection or from autoimmunity, cancer, allergy, ast graft-versus-host disease, eczema, haemophilia, thrombosis, anti-inflammatory diseases, nervous system disorders, and The present sequence encodes a protein of the invention.
the specification. The primer sets can be us in gene therapy. The primers are useful for particularly full-length cDNAs. The primers
                                                                                                                                                                                                                                      Claim
                                                                                                                                                                                                                                                                   Primer sets for synthesizing polynucleotides, particularly the 5602 full-length cDNAs defined in the specification, and for the detection and/or diagnosis of the abnormality of the proteins encoded by the full-length cDNAs -
                                                                                                                                                                                                                                                                                                                                                                                     Ota T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  28-JUL-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               07-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EP1074617-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human cDNA sequence SEQ ID NO:13526
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         26-JUN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAH15352;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                full-length cDNAs
                                                                                                                                                                                                                                                                                                                                                                                                                (HELI-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                                                   Isogai T,
                                                                                                                                                                                                                                                                                                                                                                                                                  HELIX RES
                                                                                                                                                                                                                                      SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2038 BP;
                                                                                                                                                                                                                                                                                                                                                                   Sugiyama T,
                                                                                                                                                                                                                                                                                                                                                                                                                                            2000JP-0118776.
2000JP-0183767.
2000JP-0241899.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                      ID 13526;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2000EP-0116126
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               detection; diagnosis; antisense therapy; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         99JP-0248036.
99JP-0300253.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           658
                                                                                                                                                                                                                                                                                                                                                                                                                  INST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            immune deficiencies resulting from bacterial, r from autoimmunity, cancer, allergy, asthma,
                                                                                                                                                                                                                                                                                                                                                                                  Nishikawa T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                78.2%;
86.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2
                                                                                                                                                                                                                                                                                                                                                                      Wakamatsu
                                                                                                                                                                                                                                  2537pp +
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1764
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          400 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1950
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 17.2;
Pred. No. 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ВP
                                                                                                                                                                                                                                      G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          503 G;
                                                                                                                                                                                                                                                                                                                                                                      P
                                                                                                                                                                                                                                                                                                                                                                                   Hayashi K,
                                                                                                                                                                                                                                     ROM;
                                                                                                                                                                                                                                                                                                                                                                    Nagai K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           477
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  38;
                                                                                                                                                                                                                                   English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DВ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                    Saito K,
(, Otsuki
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               22;
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                                                                                                                                                                                                                                                                                                                                                                       ..
                                                                                                                                                                                                                                                                                                                                                                                   Yamamoto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2038;
                                                                                                                                  a combination
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    infection
                                                                                                                                                                                                                                                                            detection by the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                   J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SS
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RESULT 12
AAT90543/c
ID AAT905
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Qy
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Best Local S
Matches 18
male fertility. It was obtained received to Ms41-A DNA, isolation or a sequences that showed homology to Ms41-A DNA, isolation or a sequences that showed homology to Ms41-A DNA, isolate 2m41-A genes partial cDNA clone, and use of this clone to isolate 2m41-A genes partial cDNA clone, and use of this clone to isolate 2m41-A genes 231 (AAT90542), Z33 (AAT90543) and Z35 (AAT90544) from genomic lambda libraries. The genes show a high level of conservation. Z35 may
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            detection and/or diagnosis of the abnormality of the proteins encoded by the full-length cDNAs. The primers allow obtaining of the full-length cDNAs easily without any specialised methods. AAH03166 to AAH13628 and AAH13633 to AAH18742 represent human cDNA sequences; AAB92446 to AAB95893 represent human amino acid sequences; and AAH13629 to AAH13632 represent oligonucleotides, all of which are used in the exemplification
                                                                                                                                                                                                                                                                                                                                                             exon
                                                                                                                                                                                                                                                                                                                                                                                                            exon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Key
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAT90543;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 1764 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      of the present invention.
                                                                               orthologue
                                                                                             This
                                                                                                                   Claim
                                                                                                                                          Nucleic acid encoding proteins - used to control fertility and
                                                                                                                                                                                                                                                                                                 03-JUL-1997
                                                                                                                                                                                                                                                                                                                        WO9723618-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                          exon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Zea mays
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ribozyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Zm41-A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Maize male
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAT90543
                                                                                                                                                                                                                           (GENE-) GENE SHEARS PTY LTD.
                                                                                                                                                                                                                                                                          20-DEC-1996;
                                                                                                                                                                                                                                                                                                                                                                                    intron
                                                                                                                                                                                                                                                                                                                                                                                                                                  intron
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                                                                                             DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   l Similarity
18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               male
                                                                    sequence comprises the maize Z33 Zm41-A gene. This ue of Arabidopsis Ms41-A (see AAT90522), a gene that tility. It was obtained following a database search
                                                                                                                  Fig
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 male sterile; hybrid
                                                                                                                                                                                                   Garcia D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  fertility gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                  15; 85pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                         96WO-GB03191.
                                                                                                                                                                                                                                                                                                                                                                                                        /*tag= b
936..1090
                                                                                                                                                                                                                                                                                                                                                                                                                                /*tag= a
916..935
                                                                                                                                                                                                                                                                                                                                                           /*tag= d
1168..1332
                                                                                                                                                                                                                                                                                                                                                                                  /*tag= c
1091..1167
                                                                                                                                                                                                                                                                                                                                                                                                                                                          823..915
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             511 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               76.48;
90.08;
                                                                                                                                                                                                    Hodge
                                                                                                                  English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             347 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       764
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Z33 Zm41-A.
                                                                                                                                          and
                                                                                                                                                                                                    R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 16.8;
                                                                                                                                          involved in male fertility in plants d for production of hybrid seed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  seed; transgenic
Ms41-A; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             392 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                    Perez
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                58;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DВ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ŧ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               plant; antisense;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
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                                                                     is an
confers
for
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RESULT 13
AAC56059
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         be derived from 231 via genetic rearrangements, deletions and/or insertions. 233 has subsequent deletions from 235 and is truncated, having only exons 3,5 and 6. The Ms41-A and ZM41-A genes, antisense or ribozyme sequences can be used to produce transgenic plants with controlled male fertility. Male sterile plants are useful for hybrid seed production, particularly in plants where restoration of fertility is not needed, e.g. Brassicaceae, lettuce, spinach and onions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 2582 BP; 670 A; 512 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Plant; transcription factor; gene expression; eucalyptus; pine; acaci poplar; sweetgum; teak; mahogany; bZIP; G-box binding factor; basic helix loop-helix zipper; homeotic; homeodomain; homeobox; MADS; homeodomain zipper; LIM domain; AP2; EREBs; zinc finger domain; type 2 Cys2His2; CCAAT box element; MYB; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAC56059 standard; DNA; 2444 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO200053724-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eucalyptus grandis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eucalyptus grandis transcription factor DNA sequence #190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25-JAN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAC56059
 be used to produce a plant having modified gene expression such as a be used to produce a plant having modified gene expression such as a be used to produce a plant having a cacia, poplar, sweetgum, teak, or woody plant e.g. a eucalyptus, pine, acacia, poplar, sweetgum, teak, or mahogany species or to modify the activity of a polypeptide in a plant. The transcription factors of the present invention are members from the following families of regulatory proteins: bSZIP, bZIP family of G-box binding factors, basic helix-loop-helix zipper, homeotic/homeodomain/homeobox/MADS, homeodomain zipper, LIM domain, AP2 and EREBS, zinc finger domains of type 2 Cys2His2, CCAAT box elements
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            09-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              14-SEP-2000
                                                                                                                                  The present invention relates to novel plant transcription factors from Eucalyptus grandis or Pinus radiata. The present sequence is the coding sequence for one such transcription factor. The transcription factor may
                                                                                                                                                                                                                                                           New isolated polynucle \thetatide encoding a plant transcription factor for producing a plant e.g. a^iwoody plant, preferably eucalyptus or pine, having modified gene expression or modified activity of a polypeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                          11-MAR-1999;
18-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                    (GENE-) GENESIS RES & DEV CORP LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  268 TGTTGAATCACAAGATCTGA 249
                                                                                                                                                                                                           Claim 1; Pages 94-95; 747pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 tgttggatcacaagatttga
                                                                                                                                                                                                                                                                                                                                  2000-579369/54
                                                                                                                                                                                                                                                                                                                                                                    M, McGrath A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         18;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2000WO-US06112
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99US-0149485.
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90.08;
                                                                                                                                                                                                                                                                                                                                                                        Shenk MA,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 537 G; 863 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                          Glenn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16.8;
No. 60;
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                                                                          AAH32943 to AAH37195 and AAG73514 to AAG77788 represent human colon cancer-associated nucleic acid molecules (N) and proteins (P), where the proteins are collectively known as colon cancer antigens. The colon cancer antigens have cytostatic activity and can be used in gene colors and treatment of diseases associated with inappropriate P colors cancer state with decreased expression by rectifying mutations or deletions associated with decreased expression by rectifying mutations or deletions can patient's genome that affect the activity of P by expressing colorive proteins or to supplement the patients own production of P. Colorative proteins or to supplement the patients own production of P. Colorative proteins or to supplement the patients own production of P. Colorative proteins or and P can be used in the prevention, diagnosis to express the proteins. N and P can be used in the prevention, diagnosis and treatment of coloractal carcinomas and cancers. AAH37196 to AAH37204 and AAB77789 represent sequences used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human colon cancer antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 03-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAH33702;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAH33702 standard; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              colorectal carcinoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200122920-A2
                                                                                                                                                                                                                                                                                                                                                                                                            Ruben SM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                29-SEP-1999;
03-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       28-SEP-2000; 2000WO-US26524
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo
                                                                                                                                                                                                                                                                                                                          Nucleic acids encoding 4277 human colon cancer-associated polypeptides, useful for preventing, diagnosing and/or treating colorectal cancers -
                                                                                                                                                                                                                                                                                                                                                                                                                                    (HUMA-) HUMAN GENOME SCI INC
                           present invention.

N.B. Pages 666 to 682 and page 7053 of missing at time of publication, meaning seq ID NO:1027 to 1052, 7921 and 7922.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 tgttggatcacaagattt
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                                                                                                                                                                                                                                                                                                                                                                                2001-235357/24.
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17; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             colon cancer; colon
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                                                                                                                                                                                                                                                                                                                                                                                                             Barash SC,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              99US-0157137
99US-0163280
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                643 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          74.5%;
94.4%;
                                                                                                                                                                                                                                                                                                    9803pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         335
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cancer antigen; diagnosis; detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        encoding cDNA SEQ ID
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Pred. No. 94;
                                                                                                                                                                                                                                                                                                                                                                                                             CE,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                 meaning
nd 7922.
                                                                                                                                                                                                                                                                                                                                                                                                                Rosen
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                                              the sequence I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DВ
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Sequence 335

BP;

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60 G; 105

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RESULT 15
AAV84598/c
ID AAV84598
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             18-DEC-1997
06-JUN-1997
05-SEP-1997
05-JUN-1997
06-JUN-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; secreted protein; fusion protein; gene therapy; protein therapy; diagnosis; tissue; cancer; tumour; neurodegenerative disorder; leukaemia; developmental abnormality; foetal deficiency; blood; allergy; renal; ds; immune system; asthma; lymphocytic disease; brain; hepatic; lymphoma; inflammation; ischaemic shock; Alzheimer's disease; restenosis; AIDS; cognitive disorder; schizophrenia; prostate; obesity; osteoclast; thymus; osteoprosis; arthritis; testis; lung; thyroiditis; thyroid; digestion; endocrine; metabolism; regulation; malabsorption; gastritis; neoplasm.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         04-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        10-DEC-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human secreted protein gene 188
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-MAR-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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970S-0048895.
970S-004899.
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970S-0048972.
970S-0049020.
970S-0057644.
970S-0057661.
970S-0057764.
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970S-004887.
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85.7%;
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Pred. No. 96;
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 CC rhe invention relates to nucleic acid sequences (AAV84411 to AAV84633)
CC encoding human secreted proteins (AAV88534 to AAW88756). The secreted
CC protein gene sequences are deposited with the ATCC under deposit numbers
CC ATCC 97979, 97974, 97975, 97976, 97977, 209007, 209008, 209009, 209010,
CC 209011, 209080, 209081, 209082, 209083, 209084, 209085, 209511. Host
CC cells comprising recombinant vectors containing the nucleic acid
CC sequences are used for the recombinant production of the secreted
CC proteins. The polynucleotide and amino acid sequences are useful for are
CC useful for preventing, treating or ameliorating medical conditions e.g.
CC by protein or gene therapy. Pathological conditions can be also
CC diagnosed by determining the amount of the new polypuptides in a sample
CC or by determining the presence of mutations in the new polynucleotides.
CC Specific uses are described for each of the polynucleotides, based on
CC which tissues they are most highly expressed in, and include developing
CC disorders, developmental abnormalities and foetal deficiencies, blood
CC disorders, tumours, leukemias, diseases of the immune system, autoimmune
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Li Y, Moore P
Shi Y, Soppet
                                                                                                                                                                                                                                                                                                                                                        New isolated human genes and the secreted polypeptides they encode useful for diagnosis and treatment of e.g. cancers, neurological disorders, immune diseases, inflammation or blood disorders
diseases, hepatic and renal disease, lymphomas, inflammation, aller ischemic shock, Alzheimer's and cognitive disorders, schizophrenia
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Moore PA, Ni J, Olsen HS,
Soppet DR, Wei Y, Young P
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Fischer CL, Florence C;
J, Kyaw H, Lafleur DW;
sen HS, Rosen CA, Ruben SM;
Young P, Yu G, Zeng Z;
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PRIOR APPLICATION NUMBER: U

PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO
FILING DATE:
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 2582 base paing
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; MOLECULE TYPE:
US-09-102-528-31
GENERAL INFORMATION:
APPLICANT: Gengenbac
APPLICANT: Wyse, Don
APPLICANT: Gronwald,
APPLICANT: Egli, Mar
APPLICANT: Lutz, Shi
TITLE OF INVENTION:
                                                                                                                  Sequence 1, Application US/08014326 Patent No. 5498544
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GENERAL INFORMATION:
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APPLICATION NUMBER: US/09/102,52
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy of COMPUTER: IBM PC com
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TITLE OF INVENTION:
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COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
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TITLE OF INVENTION: METHODS AND AN ACETYL COA CARBOXYLASE
TITLE OF INVENTION: FOR CONFERRING HERBICIDE TOLERANCE ANI
TITLE OF INVENTION: OIL CONTENT OF PLANTS
NUMBER OF SEQUENCES: 11
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30 (EPO)
CURRENT APPLICATION NUMBER: US/08/417,089
INFORMATION FOR SEQ ID NO: 1:
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Best Local Similarity
Matches 18; Conserv
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FILING DATE: 18-UN-1990
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/269,5
FILING DATE: 10-WOV-1998
ATTORNEY,AGENT INFORMATION:
NAME: KOWALCHY, Katherine M.
REGISTRATION NUMBER: 56,848
REFERENCE/DOCKET NUMBER: 56,848
REFERENCE/DOCKET NUMBER: 56,02.5
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
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(NFORMATION FOR SEQ ID NO: 1:
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LENGTH: 2000 base pairs
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARB: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/014,326
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MEDIUM TYPE: Floppy
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  IMMEDIATE SOURCE: 2 kb fragment of lambda clone #15-14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 21-JUL-1992 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                        APPLICANT:
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APPLICATION NUMBER: US 07/917,462
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CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                   42 GTTGGATGACAAGAGTTGTTA 62
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3100 No. 5498544west Center
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                                                                                                                                                                                          METHODS AND AN ACETYL COA CARBOXYLASE GENE FOR CONFERRING HERBICIDE TOLERANCE AND AN ALTERATION IN OIL CONTENT OF PLANTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   73.68; Score 16.2;
85.78; Pred. No. 27;
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7-1988
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Best Local Similarity
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                                                                                         INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: nucleic acid
STRANDEDNESS: single
TOOLCGY: linear
MOLECULE
                                                                                                                                                                                                                          APPLICATION NUMBER: 08/417089
FILLING DATE: 05-APR-1995
APPLICATION NUMBER: 08/014326
FILLING DATE: 05-FEB-1993
APPLICATION NUMBER: 07/917462
FILLING DATE: 21-JUL-1992
APPLICATION NUMBER: 07/538674
FILING DATE: 18-JUN-1990
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MEDIUM TYPE: Diskett
COMPUTER: IBM Compat
COMPUTER: SYSTEM: DC
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                                                                                                                                   REGISTRATION NUMBER: 30,440
REFERENCE/DOCKET NUMBER: 600.318US3
TELECOMMUNICATION INFORMATION:
TELEPHONE: 612-373-6900
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                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 2001 base pair
                                                                                                                                                                                             FILING DATE: 18-JUN-1990
ATTORNEY/AGENT INFORMATION:
NAME: Woessner, Warren D
                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: FastSEQ Vers
CURRENT APPLICATION DATA:
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MOLECULE TYPE:
                        STRANDEDNESS:
                                           TYPE:
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           TOPOLOGY:
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                        nucleic acid
EDNESS: single
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                                                       2001 base pairs
                                                                                                                       612-339-3061
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Lutz, S. M.
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            linear
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85.7%;
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Pred. No. 27;
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RESULT 6
US-08-695-421-1
: Sequence 1, Application US/08695421
: Patent No. 6268550
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US-08-930-285-1
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Best Local S
Matches 18
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Patent No. 6222099
GENERAL INFORMATION:
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GENERAL INFORMATION:
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/
FILING DATE: 13-APR-1998
CLASSIFICATION: 800
PRIOR APPLICATION DATA:
APPLICATION UMBER: PCT/US
FILING DATE: 04-APR-1996
ATTORNEY/AGENT INFORMATION:
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SEQUENCE CHARACTERISTICS:
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CORRESPONDENCE ADDRESS:
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ORIGINAL SOURCE:
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HYPOTHETICAL: NO
ANTI-SENSE: NO
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nes 18; Conserv
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85.7%;
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Pred. No. 27;
0; Mismatches
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Pred. No. 27
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RESULT 7
US-08-244-537-1
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ZIP: 55402
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKette
COMPUTER: IEM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/695,47
FILING DATE: 23-AUG-1996
CLASSIFICATION: 800
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/417089
                            GENERAL INFORMATION:
APPLICANT: 'ASHTON, ANTHONY R.
APPLICANT: JENKINS, COLIN L.D.
APPLICANT: WHITFELD, PAUL R.
TITLE OF INVENTION: MAIZE ACETYL
TITLE OF INVENTION: DNA CLONES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: 08/417089
FILING DATE: 05-APR-1995
APPLICATION NUMBER: 08/014326
FILING DATE: 05-FEB-1993
APPLICATION NUMBER: 07/917462
FILING DATE: 21-JUL-1992
APPLICATION NUMBER: 07/538674
FILING DATE: 18-JUN-1990
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ATTORNEY/AGENT INFORMATION:
NAME: Woessner, Warren D
REGISTRATION NUMBER: 30,
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CORRESPONDENCE ADDRESS:
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                 NUMBER OF SEQUENCES:
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les 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: nucleic acid
STRANDEDNESS: sing
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Somers, D. A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gengenbach,
                                                                                                                                                                                                                                                                                                                                                                                                                         linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                       single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Warren D
                                                                                                                                                                                                                                                                                                                   73.6%;
85.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       METHODS AND AN FOR CONFERRING
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                                                                                                                                                                                                                                                                                                                       Score 16.2;
Pred. No. 2
                                                                                                                                                                                                                                                                                                       Mismatches
                                                  COA CARBOXYLASE ENCODING
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COUNTRY: U.S.A. ZIP: 20005-3918

ADDRESSEE: STREET: 1

1100

CUSHMAN DARBY & CUSHMAN, L.L.P.

Washington

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; MOLECULE TYPE: DNA (genomic) US-08-244-537-1
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US-08-417-089-5
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CLASSIFICATION DATA:
APPLICATION NOMBER: GB 9125330.2
FILING DATE: 28 NOV-1991
PRIOR APPLICATION DATA:
APPLICATION DATA:
CAPPLICATION DATA:
CAPPL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 5, Application US/08417089 Patent No. 6069298 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 73.6%;
Best Local Similarity 85.7%;
Matches 18; Conservative
                                                                                                                                                                                              APPLICATION NUMBER: US.
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: ITTLE OF INVENTION: ITTLE OF INVENTION: ON UNBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE: 18-AUG-19
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
MOLECULE TYPE: -417-089-5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRANDEDNESS:
TOPOLOGY: 11
                                                                                                                                TYPE: nucleic acid
                                                                                               STRANDEDNESS:
                                                                 TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       nucleic acid
EDNESS: both
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                                DNA (genomic)
                                                                                           single
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Pred. No. 3
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US-08-695-651-5
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; MOLECULE TYPE:
US-08-695-651-5
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                                              Matches
                                                           Query Match Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/417089
FILING DATE: 05-APR-1995
APPLICATION UNMBER: 08/014326
FILING DATE: 05-FEB-1993
APPLICATION NUMBER: 07/917462
FILING DATE: 21-JUL-1992
APPLICATION NUMBER: 07/538674
FILING DATE: 18-JUN-1990
                                                                                                                                                                                 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
LENGTH: 7470 base pairs
TYPE: nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: METHOD FOR PRODUCING PLANT ACETYL COANUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 2.0
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
                                                                                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION: TELEPHONE: 612-373-6900
                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
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CITY: M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/0 FILING DATE: 12-AUG-1996 CLASSIFICATION: 435
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: Schwegman, Lundberg, Woessner & Kluth, P.A. STREET: P.O. Box 2938
                                                                                                                                                                 STRANDEDNESS: single
                                                                                                                                                                                                                                                           TELEFAX: 612-339-3061
                                                                                                                                                                                                                                                                                                                       NAME: Woessner, Warren D
REGISTRATION NUMBER: 30,4
                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER:
gttggatcacaagatttgata 22
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Lutz, S. M
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                                               Conservative
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                                                                                                                                                      linear
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85.7%; Pred. No. 32;
                                                            73.6%;
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ER: 600.318US3
                                               Score 16.2; D
Pred. No. 32;
0; Mismatches
                                               0;
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3972 GTTGGATGACAAGAGTTGTTA 3992

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; FRAGMENT TYPE:
; ORIGINAL SOURCE:
US-08-930-285-5
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                                                                                                                            Sequence 5, Patent No. (
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Best Local Similarity
Matches 18; Conserv
                                                                                                            GENERAL INFORMATION:
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LENGTH: 7470 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION: TELEPHONE: 612-339-0331
TELEFAX: 612-339-3061
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APPLICATION NUMBER: US,
FILING DATE: 13-APR-19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Regents of the University of Minnesota, et al.
TITLE OF INVENTION: TRANSGENIC PLANTS EXPRESSING ACETYL COA CARBOXYLASE GE
                                                                                                                                                                                                                        3972 GTTGGATGACAAGAGTTGTTA 3992
                                                                                                                                                                                                                                                                                                                                                                                                                HYPOTHETICAL: NO
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APPLICATION NUMBER: PCT/
FILING DATE: 04-APR-1996
ATTORNEY FACENT INFORMATION:
NAME: FACENT OF OTHER
 TITLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
                  APPLICANT:
                                 APPLICANT:
                                                 APPLICANT:
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REFERENCE/DOCKET NUMBER: 60
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: Minneapolis
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 OF INVENTION:
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                                                                                                                                           Application US/08695421
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P. O. Box 2938
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                                                               Wyse, D. L.
                                                                                                                                                                                                                                                                                        Conservative
                             Egli, M. A.
                                                Gronwa
                                                                            Somers, D. A.
                                                                                           Gengenbach,
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METHODS AND AN ACETYL COA CARBOXYLASE GENE
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Pred. No. 3:
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US-08-602-145-14/c

; Sequence 14, Application US/08602145

; Patent No. 6025336
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; MOLECULE TYPE:
US-08-695-421-5
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Best Local Similarity
                                                                                                                                                                                                                                                    GENERAL INFORMATION:
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PRIOR APPLICATION DATA:
APPLICATION UNMBER: 08/417089
FILING DATE: 05-APR-1995
APPLICATION NUMBER: 08/014326
FILING DATE: 05-FEB-1993
APPLICATION NUMBER: 07/917462
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                              3972 GTTGGATGACAAGAGTTGTTA 3992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: 18-JUN-1990
ATTORNEY/AGENT INFORMATION:
NAME: WOESSNEY, WARTEN D
REGISTRATION NUMBER: 30,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OPERATING SYSTEM: DOS
SOFTWARE: FASTSEQ Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/695,
FILING DATE: 23-AUG-1996
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                          APPLICANT: Goltry, Kristin L.

APPLICANT: Greenberger, Joel S.

TITLE OF INVENTION: DETERMINING EXPOSURE TO IONIZING RADIATION

TITLE OF INVENTION: AGENT WITH PERSISTENT BIOLOGICAL MARKERS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION: TELEPHONE: 612-373-6900
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                                                                                                                                          NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE:
STREET: P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: UZIP: 55402
                                                                                   STREET: 3000 K S
CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: 21-JUL-1992
APPLICATION NUMBER: 07/5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: Minneapolis
                                    COUNTRY: USA
ZIP: 20007-5109
                                                                        STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: 612-339-3061
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                                                                                                                          ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                            gttggatcacaagatttgata 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      nucleic acid
                                                                      D.C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            7470 base pairs
                                                                                                       E: Foley & Lardner
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Schwegman,
O. Box 2938
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FOR CONFERRING HERBICIDE TOLERANCE AND AN ALTERATION IN 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 73.6%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 16.2;
Pred. No. 32;
                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DВ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 7470;
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CURRENT APPLICATION DATA:

OPERATING SYSTEM: PC-DOS/NS-DOS SOFTWARE: Patentin Polici

Release #1.0, Version #1.30

US/08/602,145

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MOLECULE TYPE: DNA (genomic) US-08-602-145-14
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Best Local Similarity 81.8%;
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 3, Application US/08928383B Patent No. 6210921
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
LENGTH: 296 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REFERENCE/DOCKET NUMBER: 29,768
REFERENCE/DOCKET NUMBER: 076333/0112
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)672-5300
TELEFAX: (202)672-5399
TELEX: 904116
TELEFAX: (617)742-4214
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 1095 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION: NAME: Bent, Stephen A. REGISTRATION NUMBER: 29,768
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Robert W. Finberg, Jeffrey M. Bergelson,
APPLICANT: and Marshall S. Horwitz
APPLICANT: and Marshall S. Horwitz
TITLE OF INVENTION: CAR, A No. 6210921el Coxsackievirus and Adenovirus
TITLE OF INVENTION: Receptor
TITLE OF INVENTION: Receptor
NUMBER OF SEQUENCES: 26
                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE: Floppy disk
COMPOTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                               NAME: Mandragouras, Amy E. REGISTRATION NUMBER: 36,207 REFERENCE/DOCKET NUMBER: DFN-020 TELECOMMUNICATION INFORMATION: TELEPHONE: (617)227-7400 TELEPAX: (617)742-4214
                                                                                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/026,100
FILING DATE: 13-SEP-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
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| 288 TTTTGCATCATAACATTTGATA 267
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STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/08/928,383B FILING DATE: 12-SEP-1997 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 E: LAHIVE & COCKFIELD, LLP
28 State Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15-FEB-1996
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Pred. No. 42;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Mismatches
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Matches

18;

Conservative

Mismatches

0,

Gaps

0;

49;

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OTHER INFORMATION: the OTHER INFORMATION: code US-08-684-862-8
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US-08-928-383B-3
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
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Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: 07/966,040
FILING DATE: 30-DEC-1992
APPLICATION NUMBER: PCT/EP91/01361
FILING DATE: 19-JUL-1991
INFORMATION FOR SEQ ID NO: 8:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: No. 5759541el Proteins, the Preparation and Use TITLE OF INVENTION: Thereof NUMBER OF SEQUENCES: 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 5.25 inch, 360 Kb storage
MEDIUM TYPE: Diskette, 5.25 inch, 360 Kb storage
COMPUTER: IBM AT-compatible, 80286 processor
OPERATING SYSTEM: MS-DOS version 5.0
SOFTWARE: WordPerfect version 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/684,862
FILING DATE:
FILING
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/361,705
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
LENGTH: 1096 base pair
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE:
                                                                                                                                                                                                                                                                                       ORGANISM:
                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                             STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       18; .Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                           nucleic acid
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1101 Connecticut Avenue
                                                                                                                                                                                                                                                                                                                                                                         ESS: single
linear
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                                                                                                                                                                                                                                                                                          Agkistrodon rhodostoma
                                                                                                                                                                                                                                                                                                                                                     cDNA to mRNA
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         70.9%;
                                                                                                                                                                 the coding region shown in (2)(ix)(B) codes for the protein of SEQ ID NO: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 15.6;
Pred. No. 4
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      Score 15.6;
pred. No. 49
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                                                 DB 1;
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                                              Length 1096;
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COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION NUMBER: US-06/928,383B

FILING DATE: 12-SEP-1997

CLASSIFICATION:

PRIOR APPLICATION NUMBER: US-60/026,100

FILING DATE: 13-SEP-1996

ATTORNEY/AGENT INFORMATION:

NAME: Mandragouras, Amy E.

REGISTRATION NUMBER: 36,207

REFERENCE/DOCKET NUMBER: DFN-020

TELEPAX: (617)742-4214

INFORMATION: (617)742-4214

INFORMATION: GR SEQ ID NO: 1:

SEQUENCE CHARACTERISTICS:
LENGTH: 1364 base pairs

TYPE: nucleic acid

TYPE: OLDAA

MOLECULE TYPE: CDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 15
US-08-928-383B-1/c
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Search completed: December 15, 2001, 03:22:50 Job time: 7565 sec
                                                                                                8
                                                                                                                                                                                                                                                                          ; NAME/KEY: CDS
; LOCATION: 60..1157
US-08-928-383B-1
                                                                                                                                                                        Query Match
Best Local Similarity
Matches 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 1, Application US/08928383B Patent No. 6210921
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Robert W. Finberg, Jeffrey M. Bergelson,
APPLICANT: and Marshall S. Horwitz
TITLE OF INVENTION: CAR, A No. 6210921el Coxsackievirus and Adenovirus
TITLE OF INVENTION: Receptor
NUMBER OF SEQUENCES: 26
                                                                                                                                                                                                                                                                                                                                      FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
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||| | |||||||| ||| || ||
112 TGTAGTATCACAAGGTTTGCTA 91
                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE: cDNA
                                                                                          STREET: 28 St
CITY: Boston
STATE: Massac
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: USA
ZIP: 02109
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28 State Street
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Listing first 45 summaries
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                                                                      Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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SUMMARIES
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Result No.	Score 18.8	Query Match 85.5	Length DB	10 BB	ID AV532040	Description AV532040 AV532040
c 1	18.8	85.5	361	10	AV532040	AV53204
c 2	18.8	85.5	394	11	Z26559	Z26559 ATTS1651 V
C G	18.8	85.5	490	10	AB038725	AB038725
c 4	18.8	85.5	527	10	AV520789	AV520789
ი ა	18.8	85.5	530	10	AI995637	AI995637
c 6	18.4	83.6	608	11	BE977618	BE977618
c 7	18.4	83.6	934	13	CNS02BWA	AL190387
က 8	17.8	80.9	600	11	BF006444	BF006444
c 9	17.8	80.9	633	11	BF006254	BF006254
c 10	17.4	79.1	286	10	BB288727	BB288727
c 11	17.4	79.1	407	10	BE581720	BE581720
12	17.4	79.1	529	13	A0197779	A049777

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BI398487	T20362	BE118999	AV313332	AU200312	C70416	CNS04HNP	CNS029DN	CNS079F4	CNS07DUS	AZ193715	B57784	BF346432	AW963686	BE187570	AI777095	в67515	AI779714	AW034253	AQ223477	B62582	AI733664	AQ796989	W43212	AQ221715	AQ595814	AI049911	BE068225	CNS00DZ6	BE642809	BE283043	BG448853	AW695022
BI398487 952002A03	T20362 6c01g08-t7	BE118999 UI-R-CA0-	AV313332 AV313332	AU200312 AU200312	C70416 C70416 Yuji	AL291166 Tetraodon	AL187124 Tetraodon	AL435206 T7 end of	AL440954 T7 end of	AZ193715 SP_1023_B	B57784 CIT-HSP-201	BF346432 602020273	AW963686 EST375759	BE187570 EST336131	AI777095 EST258060	B67515 T22M10TR TA	AI779714 EST260593	AW034253 EST277824	AQ223477 HS_2003_B	B62582 T22F18TR TA	AI733664 an30h03.x	AQ796989 nbxb0071H	W43212 22591 Lambd	AQ221715 HS_2010_A	AQ595814 HS_2132_B	AI049911 an30h03.x	BE068225 MR4-BT036	AL077427 Drosophil	w	w	BG448853 NF003C10I	\sim

ALIGNMENTS

	source	FEATURES			COMMENT	MEDLINE	JOURNAL		TITLE	AUTHORS	REFERENCE			ORGANISM	SOURCE	KEYWORDS	VERSION	ACCESSION	DEFINITION	Locus	RESULT 1 AV532040/c
/organism="Arabidopsis thaliana" /strain="Columbia" /db_xref="columbia" /db_xref="taxon:3702" /clone="FB034f09F" /clone=11b="Arabidopsis thaliana flower buds Columbia" /tissue_type="flower buds" /note="Vector: pBluescriptII SK-; Site_1: EcoRI; Site_2: XhOI"	1361	<pre>Email: asamizu@kazusa.or.jp, URL:http://www.kazusa.or.jp/en/plant/.</pre>	Kazusa DNA Research Institute Yana 1532-3, Kisarazu, Chiba 292-0812, Japan	The First Laboratory for Plant Gene Research	Contact: Erika Asamizu	20363093	Size-Selected CDNA Libraries DNA Res. 7, 175-180 (2000)	of 12,028 non-redundant expressed sequence tags from normalized and	A large scale analysis of cDNA in Arabidopsis thaliana: Generation	Asamizu, E., Nakamura, Y., Sato, S. and Tabata, S.	1 (bases 1 to 361)	Rosidae, eurosids II; Brassicales, Brassicaceae, Arabidopsis.	Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;	Arabidopsis Chaliana	thale cress.	EST.	AV532040:1 GI:8692323		AV532040 Arabidopsis thaliana flower buds Columbia Arabidopsis thaliana cDNA clone FB034f09F 3', mRNA sequence.	361 bp mRNA EST	

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BASE COUNT
ORIGIN
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AUTHORS
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ATTS1651 Versailles-VC Arabidopsis thaliana cDNA clone VCVDH08
Similar to Ribonuclease (RNS2), mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Arabidopsis thaliana
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CNRS
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AB038725 Arabidopsis the AB
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Route de Saint-Cyr,78026 Versailles
Email: thierry@versailles.inra.fr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INRA Versailles
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thale cress. Arabidopsis thaliana Arabidopsis thaliana Arabidopsis thaliana Embryophyta; Embryophyta; Tracheophyta; Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Spermatophyta; Magnoliophyta; Enassicaceae; Arabidopsis.
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/tissue_type="whole seedlings"
/dev_stage="in vitro-grown etiolated seedlings,5
/note="Vector: pBluescript"
80 c 63 g 133 t 1 others
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/strain="ecotype Columbia"
/db_xref="taxon:3702"
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                                                                                                                                      /organism="Arabidopsis
/strain="Columbia"
/db_xref="taxon:3702"
                                                                            six-week old"
                                                                                               /clone="APZ32g06F"
/clone_lib="Arabidopsis
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Asamizu,E., Nakamura,Y., Sato,S. and Tabata,S.
A large scale analysis of cDNA in Arabidopsis thaliana: Generati of 12,028 non-redundant expressed sequence tags from normalized size-selected cDNA libraries DNA Res. 7, 175-180 (2000)
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Kazusa DNA Research Institute
Yana 1532-3, Kisarazu, Chiba 292-0812, Japan
Email: asamizu@kazusa.or.jp, URL:http://www.kazusa.or.jp/en/plant/.
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Kazusa DNA Research Institute
Yana 1532-3, Kisarazu, Chiba 292-0812, Japan
Yana 1532-1, Kisarazu, Chiba 292-0812, Japan
Email: asamizu@kazusa.or.jp, URL:http://www.kazusa.or.jp/en/plant/.
Location/Qualifiers
1. 527
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/strain="Columbia"
/strain="Columbia"
/db_xref="taxon:3702"
/clone=_lib="Arabidopsis thaliana Above-ground orgiclone_lib="Arabidopsis thaliana Above-ground orgiclone_two to six-week old plants Columbia"
/tissuc_type="aboveground organ"
/dev_stage="two to six-week old plants"
/note="Vector: pBluescriptII SK-; Site_1: EcoRI;
/tissue_type="aboveground organs"
/dev_stage="two to six-week old"
/note="Vector: pBluescriptII SK-;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
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A1995637
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Location/Qualifiers
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                                                                                                                                                     Double-stranded CDNA was blunted, ligated to SalI adaptors, digested with NotI, size-selected, and cloned into the NotI and SalI sites of the pSPORT vector."

108 c 100 g 158 t 1 others
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/tissue_type="inflorescence"
/dev_stagb="4 - 7 weeks"
                                                                                                                                                                                                                                                                                                                                                                                                       /clone="701676626"
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/cultivar="Columbia Col-0"
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Tetraodon nigroviridis genome survey sequence PUC-Ori end of clone 254C22 of library G from Tetraodon nigroviridis, genomic survey
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 bs66h08.y1 Drosophila melanogaster adult testis library Drosophila melanogaster cDNA clone bs66h08 5', mRNA sequence.
                                                       CNS02BWA
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Plate: 66 row: h column: 08
Seq primer: MIRPI reverse primer (ABI).
Location/Qualifiers
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Tissue isolation and library construction performed at the National
Institute of Diabetes and Digestive and Kidney Diseases, NIH (see
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Contact: Brian Oliver
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/strain="y[*] w[67cl]/
/db_xref="taxon:7227"
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Neptera; Endopterygota; Diptera; Brachycera;
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1 (bases 1 to 934)

1 Roest-Crollius, H., Jaillon, O., Dasilva, C., Fizames, C., Fisher, C., Roest-Crollius, H., Jaillon, O., Dasilva, C., Fizames, C., Fisher, C.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Charaterization and repeat analysis of the compact genome of the freshwater pufferfish Tetraodon nigroviridis
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Bouneau,L., Billault,A., Quetier,F., Saurin,W., Bernot,A. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              http://www.genoscope.cns.fr/Tetraodon.
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ESTs from Medicago truncatula leaves
Unpublished (2000)
Contact: Deborah A. Samac
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                                                          Fedorova, M., Pierson, B.L., Gonzales, M.B. and Ellis, L.
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/db_xref="taxon:99883"
/clone="1544C22"
/clone_ib="G"
/note="Genoscope sequence ID : COAG254BBllSP1~end
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University of Minnesota
495 Borlaug Hall, 1991 Upper Buford Circle,
Tel: 612 625 1243
Fax: 651 649 5058
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University of Minnesota name: M275482e TIGR
MTLBA56TK More information is available at:
http://chrysie.tamu.edu/medicago
Seq primer: SKmod (CTA gAA CTA gtg gAT CC).
Location/Qualifiers
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Gonzales,M.B. and Ellis,L.
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barrel medic.
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University of Minnesota
495 Borlaug_Hall, 1991 Upper Buford Circle,
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  Email: debbys@puccini.crl.umn.edu
University of Minnesota name: M275292e TIGR
MTLAU47TK More information is available at:
http://chrysie.tamu.edu/medicago
seg primer: SKmod (CTA gAA CTA gtg gAT CC).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /tissue_type="leaves and cotyledons"
/dev_stage="mixture of cotyledons from five days old plants and leaves obtained from two weeks old plants/lab_host="E. coli strain SOLR"
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/cultivar="genotype A17"
/db_xref="taxon:3880"
/clone="pbsLC-41J15"
/clone_lib="DSLC"
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Pred. No. 3.2e+02;
0; Mismatches 2
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CDNA clone pDSLC-40G21,
                                                                                                                                                                                                                                                       and cotyledons
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RESULT 10
BB288727/c
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                 P., Endo, T., Fukuda, S., Fukunishi, Y., Hara, A., Hayatsu, N., Hirozane, T., Hori, F., Ishii, Y., Ishikawa, J., Ishikawa, T., Itoh, M., Izawa, M., Kadota, K., Kagawa, I., Kai, C., Kawai, J., Kkuchi, N., Kiyosawa, H., Kojima, Y., Kondo, S., Koya, S., Kurihara, C., Kusakabe, M., Matsuyama, T., Miki, R., Mizuno, Y., Nakamura, M., Oda, H., Okazaki, Y., Ono, T., Owa, C., Saito, H., Sakai, C., Sato, K., Shibata, Y., Shigemoto, Y., Shinagawa, A., Shiraki, T., Sogabe, Y., Sugahara, Y., Suzuki, H., Tagawa, A., Takahashi, F., Tominaga, N., Toya, T., Tsunoda, Y., Watahiki, A., Watanabe, S., Yamamura, T., Yamanaka, I., Yano, R., Yasunishi, A., Yokota, T., Yoshida, K., Yoshiki, A., Yoshino, M., Muramatsu, M., and Hayashizaki, Y., RIKEN Mouse ESTS (Konno, H., et al.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BB288727 286 bp mknw
BB288727 RIKEN full-length enriched, 2 cells
clone B0200032C12 3', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 19;
                       Email: genome-res@gsc.riken.go.jp,
URL:http://genome.gsc.riken.go.jp/
Carninci,P., Nishlyama,Y., Westover,A., Itoh,M., Nagaoka,S., Sasaki,N., Okazaki,Y., Muramatsu,M. and Hayashizaki,Y.
Thermostabilization and thermoactivation of thermolabile enzymes by
                                                                                                                                                                                                                            Contact: Yoshihide Hayashizaki
Laboratory for Genome Exploration Research Group, RIKEN Genomic
Sciences Center(GSC), Yokohama Institute
The Institute of Physical and Chemical Research (RIKEN)
1-7-22 Suehiro-cho, Tsurumi-ku, Yokohama, Kanagawa 230-0045, Japan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
trehalose and its application for the synthesis of full length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Konno, H.,
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/clone_lib="postc"
/clone_lib="postc"
/tissue_type="leaves and cotyledons from five days old
/dev_stage="mixture of cotyledons from five days old
/dev_stage="mixture of cotyledons from two weeks old plants"
/lab host="E. coli strain SOLR"
/lab host="E. coli strain SOLR"
/note="Vector: pBluescript SK +/-; Site_1: EcoRI; Site_2:
XhoI; cDNA was prepared from polyA+ enriched RNA from the
mixture of cotyledons of five days old plants and leaves
of two weeks old plants. The cDNA was directionally
ligated into the Uni-ZAP XR vector from Stratagene and
packaged using Gigapack III Gold packaging extracts.
Plasmids containing cDNA inserts were excised from the
recombinant lambda-ZAP phage using Ex-Assist helper phage
and propagated in SOLR cells."
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1 to 286)
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/db_xref="taxon:3880"
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90.5%;
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Pred. No. 3.
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3.2e+02;
ches 2;
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COMMENT
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||||||||||||||
                                         The Washington Univ. Nematode EST Unpublished (1999) Contact: McCarter JP
                                                                                                                                        l (bases 1 to 407)

l (bases 1 to 407)

McCarter, J., Clifton, S., Chiapelli, B., Pape, D., Martin, J., Wylie, T., Dante, M., Marra, M., Hillier, L., Kucaba, T., Theising, B., Bowers, Y., Gibbons, M., Ritter, E., Bennett, J., Franklin, C., Tsagareishvili, R., Ronko, I., Kennedy, S., Maguire, L., Beck, C., Underwood, K., Steptoe, M., Allen, M., Person, B., Swaller, T., Harvey, N., Schurk, R., Kohn, S., Shin, T., Jackson, Y., Cardenas, M., McCann, R., Waterston, R. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BE581720 407 bp mrNA EST 09-MAY-2001 kq51c07.y1 TBN95TM-SSR Strongyloides stercoralis cDNA 5' similar WP:T19B10.2 CE16413 ;, mrNA sequence.
BE581720
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18; Conserv
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                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Nematoda; Chromadorea; Rhabditida; Panagrolaimoidea; Strongyloididae; Strongyloides.
                                                                                                                                                                                                                                                                                                                                                                              Strongyloides stercoralis
                       The Washington Univ. Nematode EST Project, 1999
                                                                                                                                                                                                                                                                                                                                                                                                     Strongyloides stercoralis.
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Washington University School of Medicine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     further details
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High-efficiency full-length cDNA cloning. Methods Enzymol. 303,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Please visit our web site (http://genome.rtc.riken.go.jp) for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      117
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    was cleaved with XhoI and BamHI. Vector: a modified pBluescript KS(+) after bulk excision from Lambda FLC I. a 66\ c 20\ g 83\ t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note-"Site_1: SalI; Site_2: BamHI; cDNA library was prepared and sequenced in Mouse Genome Encyclopedia Project of Genome Exploration Research Group in Riken Genomic Sciences Center and Genome Science Laboratory in RIKEN. Division of Experimental Animal Research in Riken contributed to prepare mouse tissues. 1st strand cDNA was
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /clone="B020032C12"
/clone_lib="RIKEN full-length enriched,
/tissue_type="egg"
/dev_stage="2_cells"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /organism="Mus musculus"
/db_xref="taxon:10090"
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94.7%;
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Pred. No. 4.3e+02;
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les 18; Conservative
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High Throughput Sequencing Center
University of Washington
401 Queen Anne Avenue North, Seattle, WA 98109, USA
401 Queen Anne Avenue North, Seattle, WA 98109, USA
Tel: (206) 616-3618
Fax: (206) 616-3887
Email: jwallace@u.washington.edu
Email: jwallace@u.washington.edu
Clones are derived from the human BAC library RPCI-11. For BAC
Clones are derived from the human BAC library RPCI-11. For BAC
pleter@dejong.med.buffalo.edu). Clones may be purchased from
(pleter@dejong.med.buffalo.edu). Clones may be purchased from
BACPAC Resources (http://bacpac.med.buffalo.edu/ordering_bac.htm)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TTGATCACAAGATTTGATA 316
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The library was constructed by Dr. Thomas Nutman
NIAID, NIH (tnutmann@nih.gov). DNA Sequencing by:
University Genome Sequencing Center St. Louis.
High quality sequence stop: 384.
Light quality sequence stop: 184.
1. 407
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Fax: 314 286 1810
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 529)
Mahairas,G.G., Wallace,J.C., Smith,K., Swartzell,S., Holzman,T., Keller,A., Shaker,R., Furlong,J., Young,J., Zhao,S., Adams,M.D. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence-tagged connectors: A sequence approach to mapping scanning the human genome
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AQ497779.1 GI:4697902
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact: Mahairas GG, Wallace JC, Hood
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                152
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//lab_host="XL-1 Blue MRF' (Stratagene)"
//lab_host="XL-1 Blue MRF' (Stratagene); Site_1:
//note="Vector: Lambda Uni-ZAP XR (Stratagene); Solated from gerbils
rhabditiform larvae which had been isolated from gerbils
from experimentally infected dogs. cDNA was constructed
from experimentally infected dogs. cDNA was constructed
and, using adaptors, was cloned unidirectionally into the
vector from the EcoRI site to the XhoI site. The library
has an unamplified titer of 1 x 10E5 pfu/ml and an
amplified, undiluted titer of 9 x 10E11 pfu/ml. The
average insert size of the unamplified library is 675 bp
(range, 100-1700)."
18 t
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/db_xref="taxon:6248"
/clone_lib="TBN95TM-SSR"
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94.7%;
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Pred. No. 4.6e+02;
0; Mismatches 1;
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High quality sequence stop: 529
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Class: BAC end
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AW695022
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EST.
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He,X.-Z., Shadle,G., Scott,A.D., Harris,A.R., Gonzales,R.A., Be.,
C.J., Flores,H.R., Inman,J.T., Weller,J.W., May,G.D. and Dixon,R.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Medicago truncatula
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
Rosidae; eurosids I; Fabales; Fabaceae; Papillonoideae; Trifolieae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unpublished (2000) on Apr 14, 2000 this sequence version replaced gi:7569784. Contact: Dixon RA
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                                                                                                                                                                                                                                                                                                                                                                                                                             Email: radixon@noble.org
Insert Length: 674 Std Error: 0.00
Plate: 082 row: E column: 04
Seg primer: TCACACAGGAAACAGCTATGAC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2510 Sam Noble Parkway, Ardmore,
Tel: 580 221 7302
Fax: 580 221 7380
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The Samuel Roberts Noble Foundation Roberts Noble Foundation
The Samuel Roberts Noble Foundation Rober
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/db_xref="taxon:9606"
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/clone_lib="RPCI-11 Human Male BAC Library"
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                                             /tissue_type="stem"
/dev_stage="Pooled developmental"
/note="vector: Lambda Zap; Contains a
internodal stem segments"
142 c 129 g 195 t
                                                                                                                                                                                                                                                     /organism="Medicago truncatula"
/db_xref="taxon:3880"
/clone="NF082E0457"
                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
1. .646
                                                                                                                                                                                                                  /clone_lib="Developing stem"
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Pred. No. 4.8e+02;
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mRNA sequence.
BE283043
BE283043.1 GI:9159023
                                                     BE283043 762 bp mRNA
601101323F1 NCI_CGAP_Lu29 Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           217 Plant Science Building, Fayetteville, AR 72701, USA Tel: 501 575 5191
Fax: 501 575 7601
Email: kkorth@comp.uark.edu
Insert Length: 671 Std Error: 0.00
Plate: 003 row: C column: 10
Seq primer: TCACACAGGAAACAGCTATGAC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Korth K
Dept. of Plant Pathology
University of Arkansas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 (bases 1 to 671)

KOrth,K., Scott,A.D., Harris,A.R., Gonzales,R.A., Bell,C.J., Flo., H.R., Inman,J.T., Weller,J.W. and May,G.D.

Expressed Sequence Tags from the Samuel Roberts Noble Foundation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Medicago truncatula
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
Rosidae; eurosids I; Fabales; Fabaceae; Papilionoideae; Trifolieae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BG448853 671 bp mRNA EST 16-MAR-2001 NF003C10IN1F1082 Insect herbivory Medicago truncatula cDNA clone NF003C10IN 5', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BG448853.1 GI:13367634
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Medicago truncatula insect herbivory library
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                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                          /clone_lib="Insect herbivory"
/tissue_type="local and systemic leaves"
/dev_stage="mature"
/note="Vector: Lambda 2ap; Library was produced from fully
/note="Vector: Lambda 2ap; Library was produced from fully
expanded M. truncatula leaves of plants fed upon by
spodoptera exigua (beet armyworm) for 24 hours. Systemic
(undamaged leaves from injured plants) and wounded leaves
were harvested and pooled."

151 c 131 g 203 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /organism="Medicago truncatula"
/db_xref="taxon:3880"
/clone="NF003C10IN"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            .671
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94.7%;
                                                                                                                                                                                                                                              79.1%; Score 17.4; DB 94.7%; Pred. No. 5e+02; Live ;0; Mismatches
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Pred. No. 4.9e+Q2;
0; Mismatches 1;
                                                         RNA EST
Mus musculus cDNA clone
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IMAGE:3493829 5',
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AUTHORS
TITLE
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Search completed: December 15, Job time: 4989 sec
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Best Local S
Matches 18
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                                                                                                692 TGTTGGATCACAAGACTTG 710
                                                                                                                   tgttggatcacaagatttg 19
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                                                                                                                                                                          l Similarity
18; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 762)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Gilbert Smith,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mus musculus
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                                                                                                                                                                                                                                                                                       /note="Organ: lung; Vector: pCMV-SPORT6; Site_1: SalI; Site_2: NotI; Cloned unidirectionally. Primer: Oligo of Library constructed by Life Technologies. Investigator providing samples: Gilbert Smith, NIH"

a 172 c 207 g 190 t
                                                                                                                                                                                                                                                                                                                                                                                       /strain="C2ECH II (feral)"
/db_xref="taxon:10090"
/clone="IMAEE:3493829"
/clone_lib="NCI_CGAP_Lu29"
/tissue_type="spontaneous tumor, metastatic stem cell origin."
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /organism="Mus musculus"
                                                                                                                                                                                          79.18;
94.78;
                    2001,
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                                                                                                                                                                        Score 17.4; DB 10
Pred. No. 5.1e+02;
0; Mismatches 1
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 2000000000
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1477
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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35 36 37 38 39	25 25 30 31 32 33 34	15 17 18 19 20 21 23	7 8 9 10 11 12 13	1654221	No.
794.8 794.8 792.6 792.6 790.6 790.4 790.2 789.8		801.6 801.6 801.6 801.6 801.6 801.6 801.6 801.4 799.4	803.22 803.22 803.22	1477 814.2 814.2 810.2 810.2 810.2	Score
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13 17 16 29 12 29 29		9 13 17 17 12 12 12 17 17 23	01009970	775307	DB
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Sequence 12, Appl Sequence 1, Appli Sequence 1, Appli Sequence 1, Appli Sequence 25, Appli Sequence 3, Appli Sequence 3, Appli Sequence 1, Appli	399, 406, 711, 716, 716, 716, 13, 13, 13, 750,	160, A 11, A 11, A 21, A 22, A 28, A 79, A 7439, 7439, 7441,	Sequence 1256, Ap Sequence 1291, Ap Sequence 1, Appli Sequence 1, Appli Sequence 1, Appli Sequence 1, Appli Sequence 5, Appli Sequence 5, Appli	1, API 1, API 1, API 1, API 14, API 14, API 14, API	Description

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Sequence 1, Application US/09380826A
GENERAL INFORMATION:
APPLICANT: Chappel, Rod
TITLE OF INVENTION:
LEPTOSPIRA PATHOGENS
FILE REFERENCE: DARETS, 001APC
CURRENT APPLICATION NUMBER: US/09/380,826A
CURRENT FILING DATE: "3,999-11-22
PRIOR APPLICATION NUMBER: POT/AU98/00145
PRIOR FILING DATE: 1998-03-06
PRIOR APPLICATION NUMBER: AU PO5494/97
PRIOR FILING DATE: 1997-03-07
NUMBER OF SEQ ID NOS: 26
SOPTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 117
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 1
US-09-380-826A-1
                                                                                                                                                                                                                                                                                                                                                     Query Ma
Best Loc
Matches
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TYPE: DNA
ORGANISM: Leptospira
-09-380-826A-1
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                                                            ctttccgaaaggaaagctaataccggatagtcctgttggatcacaagatttgataggtaa
                         ccaaggcgacgatcggtagccggcctgagagggtgtccggccacaatggaactgagacac
cggatttgtaagtcaggtgtgaaaactgcggggctcaacccgtggcctgcacttgaaacta
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US-09-726-774-9
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Pred. No. 0;
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US-08-642-229-1

Sequence 1, Application US/08642229

GENERAL INFORMATION:
APPLICANT: Herwig, Russell P.
APPLICANT: Bielefeldt, Angela R.
APPLICANT: Stensel, H. David
APPLICANT: Strand, Stuart E.
APPLICANT: STRAND, Degradation
TITLE OF INVENTION: Filamentous
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   T: Stensel, H. David
T: Strand, Stuart E.
INVENTION: Degradation
INVENTION: Filamentous
                                                                 US/08642229
     of Enviror
Bacterium
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US-08-642-229-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Sheiness, Diana K.

REGISTRATION NUMBER: 35,356

REFERENCE/DOCKET NUMBER: UOFW19233

TELECOMMUNICATION INFORMATION:

TELEPHONE: (206) 682 8100

TELEPAX: (206) 224 0779

TELEFAX: (206) 224 0779

TELEFAX: (206) 224 0779

TELEX: 4938023

INFORMATION FOR SEQ ID NO: 1:

SEQUENCE CHARACTERISTICS:
LENGTH: 1536 base pairs
                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
Matches 1113; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/642,229
FILING DATE:
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                   STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: nucleic acid
DESCRIPTION: "16S ribosomal
HYPOTHETICAL: NO
ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
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APPLICATION NUMBER: 1
FILING DATE: 20-MAY-
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STREET: 1420 Fifth.
CITY: Seattle
STATE: Washington
COUNTRY: USA
ZIP: WA 98101-2347
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: nucleic acid
                                                                               tgataggtaaagatttattgcttggagatgagcccgcggccgattagctagttggtgagg
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aaaccctgaagcagcgacgccgcgtgaacgaaggaaggtcttcggattgtaaagttcatta
                                                actgagacacggtccatactcctacgggaggcagcagttaagaatcttgctcaatggggg
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                                    ACTGAGACACGGCCCAGACTCCTACGGGAGGCAGCAGTGGGGAATTTTGGACAATGGGCG
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Pred. No. 3.5e
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US-08-965-623-1
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                                                                                                                                                                                                                   Query Match
Best Local
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TELEX: 4938023
INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: n
DESCRIPTION: "
HYPOTHETICAL: NO
ANTI-SENSE: NO
ORIGINAL SOURCE:
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APPLICANT: S
APPLICANT: S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION US/08/965,623
FILING DATE: November 6, 1997
CLASSIFICATION: 435
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REGISTRATION NUMBER: 35,356
REFERENCE/DOCKET NUMBER: UOFW111488
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 682 8100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 30-APRIL-19: ATTORNEY/AGENT INFORMATION: NAME: Sheiness, Diana K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/246,865
FILING DATE: 20-MAY-1994
APPLICATION NUMBER: US 08/599,867
FILING DATE: 12-FEBRUARY-1996
APPLICATION NUMBER: US 08/642,229
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STRANDEDNESS: single
TOPOLOGY: linear
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ZIP: WA 98101-2347
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1420 Fifth Avenue, Suite 2800
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Stensel, H. David .
Strand, Stuart E. .
Herwig, Russell P.
VENTION: Degradation of Environmental
VENTION: Filamentous Bacterium
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"16S ribosomal DNA"
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APPLICANT: Andersen, Lene Nonboe
APPLICANT: Schulein, Martin
APPLICANT: Schulein, Martin
APPLICANT: Hogernvad, Mads Eskelund
APPLICANT: Bjoernvad, Mads Eskelund
APPLICANT: Glad, Sanne O. Schroeder
APPLICANT: Glad, Sanne O. Schroeder
APPLICANT: Filler Soeren
APPLICANT: Soeren
APPLICATION: NOVEL Pectate Lyases
FILE REFERENCE: 5378.200-US
CURRENT APPLICATION NUMBER: US/09/198,955
CURRENT FILLING DATE: 1998-11-24
EARLIER APPLICATION NUMBER: 1344/97
EARLIER APPLICATION NUMBER: 1344/97
EARLIER APPLICATION NUMBER: 60/067,249
EARLIER APPLICATION NUMBER: 60/067,249
EARLIER APPLICATION NUMBER: 60/067,249
EARLIER APPLICATION NUMBER: 09/07,240
EARLIER APPLICATION NUMBER: 09/07,240
EARLIER APPLICATION NUMBER: 09/07,240
EARLIER FILING DATE: 1997-12-02
EARLIER FILING DATE: 1998-05-06
EARLIER FILING DATE: 1998-05-06
EARLIER APPLICATION NUMBER: 09/073,684
EARLIER APPLICATION NUMBER: 09/073,684
EARLIER FILING DATE: 1998-05-06
EARLIER APPLICATION NUMBER: 09/184,217
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SOFTWARE: FASESEQ for
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; LENGTH: 1508
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; ORGANIZM: Bacillus :
US-09-198-955-14
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Best Local Similarity
Matches 921; Conserv
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960 991 1020 1048 1080 720 755 780 813 840 873 600 635

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APPLICANT: Andersen, Lene Nonboe
APPLICANT: Bjornvad, Mads Eskelund
TITLE OF INVENTION: Novel Mannanases
FILE REFERENCE: 5440.204-US
CURRENT APPLICATION NUMBER: US/09/339,159A
CURRENT FILING DATE: 1999-06-24
EARLIER APPLICATION NUMBER: 60/106,054
EARLIER FILING DATE: 1998-10-28
EARLIER FILING DATE: 1998-10-28
EARLIER FILING DATE: 1998-01-28
EARLIER FILING DATE: 1999-03-09
EARLIER APPLICATION NUMBER: 60/123,543
EARLIER APPLICATION NUMBER: 60/123,623
EARLIER APPLICATION NUMBER: 60/123,623
EARLIER FILING DATE: 1999-03-10
EARLIER FILING DATE: 1999-03-10
EARLIER APPLICATION NUMBER: 60/123,641
EARLIER APPLICATION NUMBER: 60/123,641
EARLIER APPLICATION NUMBER: 60/123,641
EARLIER APPLICATION NUMBER: 60/123,642
EARLIER APPLICATION NUMBER: 60/123,642
EARLIER FILING DATE: 1999-03-10
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Best Local Similarity 61.7
Matches 921; Conservative
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ORGANISM: Bacillus
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APPLICANT: Schulein, Mar
APPLICANT: Schnorr, Kirk
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                                ccgccgtcacaccagctgagtggggagcacccgaagtggtctttgttaaccgta-agga
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FILE REFERENCE: 5378.200-US
CURRENT APPLICATION NUMBER: US/09/694,531
CURRENT FILING DATE: 2000-10.23
PRIOR APPLICATION NUMBER: 09/199,955
PRIOR FILING DATE: 1998-11-24
PRIOR APPLICATION NUMBER: 1344/97
PRIOR FILING DATE: 1997-11-24
PRIOR FILING DATE: 1997-11-24
PRIOR APPLICATION NUMBER: 1344/97
PRIOR FILING DATE: 1997-12-02
PRIOR APPLICATION NUMBER: 60/067,240
PRIOR FILING DATE: 1998-12-02
PRIOR APPLICATION NUMBER: 09/073,684
PRIOR APPLICATION NUMBER: 09/073,684
PRIOR APPLICATION NUMBER: 09/073,684
PRIOR APPLICATION NUMBER: 09/184,217
PRIOR APPLICATION NUMBER: 09/184,217
PRIOR FILING DATE: 1998-11-02
NUMBER OF SEQ ID NOS: 32
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 14
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US-09-694-531-14
Sequence 14, Application US/09694531
GENERAL INFORMATION:
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APPLICANT: Schulein, Marti
APPLICANT: Schulein, Marti
APPLICANT: Bjornvad, Mads
APPLICANT: Moller, Soren
APPLICANT: Glad, Sanne O.
APPLICANT: Kauppinen, Mark
APPLICANT: Schnorr, Kirk
                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 1508
TYPE: RNA
ORGANISM: Bacillus:
S-09-694-531-14
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Best Local Similarity
Matches 921; Conserv
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APPLICANT: Kongsbak, Lars
TITLE OF INVENTION: Novel Pectate
FILE REFERENCE: 5378.200-US
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Kauppinen, Markus S.
Schnorr, Kirk
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Lange, Niels E.
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Moller, Soren
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, 61.7% Pred. 1
cvative 220; Misr
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APPLICANT: SORCKINE, Alexel
APPLICANT: Bolotine, Alexandre
APPLICANT: Lapidus, Alla
TITLE OF INVENTION: Methods For Monitoring Mu:
TITLE OF INVENTION: Expression
FILE REFERENCE: 5963.000-US
CURRENT APPLICATION NUMBER: US/60/279,526
CURRENT FILING DATE: 2001-03-27
NUMBER OF SEQ ID NOS: 2232
SOPTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1256
LENGTH: 5643
LENGTH: 5643
CAGANISM: B. Clausii
FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)...(5643)
OTHER INFORMATION: n = A,T,C or G
US-60-279-526-1256
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Best Local Similarity 75.2
Matches 1130; Conservative
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RESULT 8
US-09-680-598A-1291/c
US-09-680-598A-1291/c
; Sequence 1291, Application U:
; GENERAL INFORMATION:
; APPLICANT: Randy M. Berka
; APPLICANT: Ib Groth Clause

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NAME/KEY: misc_feature
LOCATION: (1)...(2484)
OTHER INFORMATION: n =
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Best Local Similarity 75.48;
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TYPE: DNA
ORGANISM: Bacillus 1
FEATURE:
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Pred. No. 1.2e-247;
0; Mismatches 324;
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RESULT 9
US-09-745-476-1
US-09-745-476-1
; Sequence 1, Application US/09745476
; GEMERAL INFORMATION:
APPLICANT: CANON INC.
TITLE OF INVENTION: Preparation of Poly-hidroxyalkanoic
FILE REFERENCE: 4351008
; CURRENT APPLICATION NUMBER: US/09/745,476
; CURRENT FILING DATE: 2000-12-26
; NUMBER OF SEQ ID NOS: 1
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; SEQ ID NO 1
; LENGTH: 1501
; TYPE: DNA
; ORGANISM: Pseudomonas j
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RESULT 10
US-09-748-205-1
Sequence 1, Application US/09748205
GENERAL INFORMATION:
APPLICANT: Canon Inc.
TITLE OF INVENTION: Polyhydroxyalkanoate it
TITLE OF INVENTION: those are used for the
FILE REFERENCE: 4351009
CURRENT APPLICATION NUMBER: US/09/748,205
CURRENT FILING DATE: 2000-12-27
NUMBER OF SEQ ID NOS: 1
SEQ ID NO 1
LENGTH: 1501
                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMB
CURRENT FILING DATE: 20
NUMBER OF SEQ ID NOS: 1
SEQ ID NO 1
LENGTH: 1501
TYPE: DNA
ORGANISM: Pseudomonas J.
US-09-748-205-1
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SEQ ID NO 1
LENGTH: 1501
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APPLICANT: Canon Inc.
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CURRENT FILING DATE: 2001-02-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: Polyhydroxyalkanoate containing TITLE OF INVENTION: monomer unit, and method for purile REFERENCE: 4396021
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Pred. No. 1.2e-247;
0; Mismatches 318;
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                                     RESULT 12
US-09-793-920A-1
US-09-793-920A
; Sequence 1, Application US/09793920A
; GENERAL INFORMATION:
APPLICANT: Canon Inc.
TITLE OF INVENTION: monomer unit, and method for producing the same.
TITLE OF INVENTION: monomer unit, and method for producing the same.
FILE REFERENCE: 4396021
CURRENT APPLICATION NUMBER: US/09/793,920A
; CURRENT FILING DATE: 2001-02-28
; NUMBER OF SEQ ID NOS: 1
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APPLICANT: CANON INC.

APPLICANT: CANON INC.

TITLE OF INVENTION: Polyhydroxyalkanoate Synthase and Gene Ence
FILE REFERENCE: 4051021

CURRENT APPLICATION NUMBER: US/09/821,016

CURRENT FILING DATE: 2001-03-30

NUMBER OF SEO ID NOS: 11

SOFTWARE: Microsoft Word

SEO ID NO 5

LENGTH: 1501

TYPE: DNA

ORGANISM: Pseudomonas jessenii P161; BP-7376

FEATURE:
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Mon Dec 17 07:48:11 2001

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APPLICANT: Berka, Randy M.

APPLICANT: Clausen, Ib Groth
APPLICANT: Sorokine, Alexei
APPLICANT: Bolotine, Alexei
APPLICANT: Bolotine, Alexei
APPLICANT: Bolotine, Alexandre
APPLICANT: Lapidus, Alla
TITLE OF INVENTION: Expression
FILE REFERRNCE: 9563.000-US
CURRENT APPLICATION NUMBER: US/60/279,526
CURRENT FILLING DATE: 2001-03-27
NUMBER OF SEO ID NOS: 2232
SOFTWARE: FASTSEQ for Windows Version 4.0
SEO ID NO 5
SEO ID NO 5
SEO ID NO 5
LENGTH: 5901
TYPE: DNA
ORGANISM: B. Clausii
FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)...(5901)
OTHER INFORMATION: n = A,T,C or G
US-60-279-526-5
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GENERAL INFORMATION:
APPLICANT: Berka, Randy M.
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Best Local Similarity 75.0
Matches 1128; Conservative
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Pred. No. 2.7e-247;
0; Mismatches 330; Indels 47;
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Qy 1395 taaccgtaaggagacagactactaaggtgaaactcgtaaagggggtgaa
Oy 1335 gaccttgtacacaccgcccgtcacaccacctgagtggggagcacccgaa
Qy 1275 cccatgaagtcggaatcgctagtaatcgcggatcagcatgccgcgggtga
Qy 1215 gcaagagggagctaatctctaaaagtcggtcccagttcggattggggtc
Qy 1155 ggcctttatgtccagggccacacacgtgctacaatggccgatacagagg
Oy 1095 actggtacgaaactgccggtgacaaaccggaggaaggcggggatgacgt
Qy 1036 ttgggttaagtcccgcaacgagcgcaacccct-atcgtatgttgctacct
Qy 976 tgagccttcgggcagattcacaggtgctgcatggttgtcgtcagctcgt
Qy 919 tgataccccaaaaacctcacctgggcttgacatggatctgaatcatg
Qy 859 tgaaactcaaaggaattgacgggggtccgcacaagcggtggagcatgtgg
Qy 801ccctcagtaacgaacctaacggattaagtagaccgcctggggactatg
Oy 741 gattagataccccggtaatccacgccctaaacgttgtctaccagttgttg
Oy 681 cgaaggcgacttgctggctcaaaactgacgctgaggcacgaaagcgtggg
Qy 621 ggcaagtggaattccaggtgtagcggtgaaatgcgtagatatctggagga
Qy 561 gaaaactgcgggctcaacccgtggcctgcacttgaaactacaagtctgga
Qy 501 aagcgttgttcggaatcattgggcgtaaagggtgcgtaaggcggatttgta
Oy 452aaagcaccggctaactacgtgccagcagccgcggtaata
Db 5142 CGTAAAGCTCTGTTGTGAGGGAAGAAGCGGTACCGTTCGAATAGGGCGGT

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US-08-520-946-160
; Sequence 160, Application:
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OPERATING SYSTEM: PC-DOS/MS-DOS.
SOFTWARE: Patentin Release #1.0, Version #1
CURRENT APPLICATION DATA:
APPLICATION NATA:
APPLICATION UMBER: US/08/520,946
FILING DATE:
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: CARROLL, PETER G.
REGISTRATION NUMBER: 32,837
REFERENCE/DOCKET NUMBER: FORS-01756
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 705-8410
TELEPAX: (415) 397-8338
INFORMATION FOR SEO ID NO: 160:
SEQUENCE CHARACTERISTICS:
LENGTH: 1555 base pairs
TYPE: NUCLEIC acid
STRANDEDNESS: double
TOPOLOGY: Inhear
MOLECULE TYPE: DNA (GENOMIC)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 54.3%;
Best Local Similarity 75.3%;
Matches 1134; Conservative
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatit
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APPLICANT: LYAMICHEV, VICTOR I.
APPLICANT: OLIVE, DAVID M.
TITLE OF INVENTION: RAPID DETECTION AND
TITLE OF INVENTION: PATHOGENS
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STREET: 220 MONTGOMERY STREI
CITY: SAN FRANCISCO
STATE: CALIFORNIA
COUNTRY: UNITED STATES OF AVECTOR OF 
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Pred. No. 4.2e-247;
0; Mismatches 324;
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CURRENT FILING DATE: 2000-08-14
PRIOR APPLICATION NUMBER: JP P1998-356229
PRIOR FILING DATE: 1998-12-15
NUMBER OF SEQ ID NOS: 2
SOFTWARE: PATENTIN Ver. 2.1
SEQ ID NO 1
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RESULT 3
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Matches 1114; Conserv
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LENGTH: 1480
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ORGANISM: Unknown
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Sequence 1, Application US/09902693
GEMERAL INFORMATION:
APPLICANT: MITARA, YASUSHITO
APPLICANT: TAKEUCHI, SONOKO
APPLICANT: TONOUCHI, NOOTO
APPLICANT: TONOUCHI, NOOTO
APPLICANT: FUDOU, RYOSUKE
APPLICANT: FUDOU, RYOSUKE
APPLICANT: YOKOZEKI, KENZO
TITLE OF INVENTION: D-XYLUUOSE
FILE REFERENCE: 0010-1015-0
CURRENT APPLICATION NUMBER: US/09/902,693
CURRENT FILING DATE: 1999-07-02
PRIOR APPLICATION NUMBER: JP 10-193472
PRIOR FILING DATE: 1998-07-08
PRIOR FILING DATE: 1998-07-08
PRIOR FILING DATE: 1998-10-30
PRIOR APPLICATION NUMBER: JP 10-310398
PRIOR APPLICATION NUMBER: JP 11-12244
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; PRIOR FILING DATE: 1999-01-20; NUMBER OF SEQ ID NOS: 5
; SOFTWARE: PATENTIN VET. 2.1
; SEQ ID NO 1
; LENGTH: 1438
; TYPE: DNA
; ORGANISM: Unknown Organism
; FEATURE:
; OTHER INFORMATION: Description
; OTHER INFORMATION: N at positin
US-09-902-693-1
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US-09-897-516-423
   \mathbf{s}
                                                                                                                                APPLICANT: Corbin, David R.
APPLICANT: Goldman, Barry J.
APPLICANT: Hinkle, Gregory J.
APPLICANT: Hinkle, Joseph E.
APPLICANT: Huesing, Joseph E.
APPLICANT: Krasomil-Osterfeld, Karina C.
APPLICANT: Slater, Steven C.
APPLICANT: Spiridonov, Sergei
TITLE OF INVENTION: Xenorhabdus sp. Genome SeriLE REFERENCE: 38-21(51847)B
CURRENT APPLICATION NUMBER: US/09/897,516
CURRENT FILING DATE: 2001-06-29
PRIOR APPLICATION NUMBER: US 60/215, 161
PRIOR FILING DATE: 2000-06-30
NUMBER: OF SCO. 101 NO. 2000-06-30
                                                                                        NUMBER OF SEQ ID NOS: 8409
SEQ ID NO 423
LENGTH: 83080
                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                    Sequence 423,
NAME/KEY: CDS
LOCATION: (38740)..(39948)
-09-897-516-423
                                             TYPE: DNA
ORGANISM: Xenorhabdus
FEATURE:
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367 aaaggaattgacgggggtccgcacaagcggtggagcatgtggtttaattcgatgataccc 926 	867 1514	Qy dd
807 gtaacgaacctaacggattaagtagaccgcctggggactatgctcgcaagagtgaaactc 866 	807 1454	Qy dd
48 taccccggtaatccacgccctaaacgttgtctac-cagttgttgggggttttaaccctca 806 	748 1394	Qy Db
188 gacttgctggctcaaaactgacgctgaggcacgaaagcgtgggtagtaaacgggattaga 747 	1334	Dh Qy
28 ggaattccaggtgtagcggtgaaatgcgtagatatctggaggaacaccagtggcgaaggc 68 	628 1274	Qy Db
68 gcgggctcaacccgtggcctgcacttgaaactacaagtctggagtttgggagaggcaagt 627 	568 1214	Db Qy
108 gttcggaatcattgggcgtaaagggtgcgtaggcggatttgtaagtcaggtgtgaaaact 567 	508 1154	Оу
48 gcctaaagcaccggctaactacgtgccagcagccgcggtaatacgtatggtgcaagcgtt 50 	1094	Qy dd
03 gttcattaggcaggaaaaataagcagcaatgtgatgatggtacct 44	403 1034	Db Qy
43 aatggggaaacctgaagcagcgac 	343 974	Db Qy
183 acaatggaactgagacacggtccatactcctacgggaggcagcagttaagaatcttgctc 342 	283 914	Qy Db
23 tggtgaggtaatggctcaccaaggcgacgatcggtagccggcctgagagggtgtccggcc 282 	a t	dd Vy
63 acaagatttgataggtaaagatttattgcttggagatgagcccgcggccgattagctagt 222 	163 794	Оу
03 cctccgagtctgggataactt 36 cccgatggagggggataacca	103 736	Ωy
61 caatac	6	DP QV
actaa ttgaa	6	Db Qy
Match 51.7%; Score 763.4; DB 6; Length 83080; Local Similarity 73.4%; Pred. No. 1.9e-252; es 1100; Conservative 0; Mismatches 356; Indels 43; Gaps	Query Mat Best Loca Matches 1	

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APPLICANT: Goldman, Barry S.
APPLICANT: Hinkle, Gregory J.
APPLICANT: Huesing, Joseph E.
APPLICANT: Krasomil-Osterfeld, Karina C.
APPLICANT: Krasomil-Osterfeld, Karina C.
APPLICANT: Slater; Steven C.
APPLICANT: Slater; Steven G.
APPLICANT: Spiridonov, Sergei
TITLE OF INVENTION: Xenorhabdus sp. Genome S
FILE REFERENCE: 38-21(51847)B
CURRENT APPLICATION NUMBER: US/09/897,516
CURRENT FILING DATE: 2001-06-29
PRIOR APPLICATION NUMBER: US 60/215, 161
PRIOR FILING DATE: 2000-66-30
NUMBER OF SEG ID NOS: 8409
SEQ ID NO 3852
LENGTH: 83080
TYPE: DNA
ORGANISM: Xenorhabdus sp.
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APPLICANT: GOLDMan, Barry S, APPLICANT: GOLDMan, Barry S, APPLICANT: Hinkle, Gregory APPLICANT: Huesing, Joseph I, APPLICANT: Krasomil-Osterfel APPLICANT: Malvar, Thomas M, APPLICANT: MALVAR, MA
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US-09-897-516-3852
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Best Local Similarity
Matches 1100; Conser
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LOCATION: (1362)..(1466)
-09-897-516-3852
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                                         Score 763.4; D
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US-09-997-516-3853
; Sequence 3853, Application
; GENERAL INFORMATION:
; APPLICANT: COIDIN, David R
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Best Local Similarity 73.4
Matches 1100; Conservative
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LOCATION: (1931)..(2044)
-09-897-516-3853
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TYPE: DNA
ORGANISM: Xenorhabdus
FEATURE:
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APPLICANT: Goldman, Barry S.
APPLICANT: Hinkle, Gregory J.
APPLICANT: Huesing, Joseph E.
APPLICANT: Krasomil-Osterfeld, Karina C.
APPLICANT: Krasomil-Osterfeld, Karina C.
APPLICANT: Slater, Steven C.
APPLICANT: Slater, Steven C.
APPLICANT: Spiridonov, Sergei
TITLE OF INVENTION: Xenorhabdus sp. Genome Stelle REFERENCE: 38-21(51847)B
CURRENT APPLICATION NUMBER: US/09/897,516
CURRENT FILING DATE: 2001-06-29
PRIOR APPLICATION NUMBER: US 60/215, 161
PRIOR FILING DATE: 2000-06-30
NUMBER OF SEQ ID NOS: 8409
SEO ID NO 3854
LENGTH: 83080
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; NAME/KEY: CDS
; LOCATION: (5826)..(6755)
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US-09-897-516-3854
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APPLICANT: Goldman, Barry
APPLICANT: Hinkle, Gregor
APPLICANT: Huesing, Josep
APPLICANT: Krasomil-Oster
APPLICANT: Malvar, Thomas
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Best Local Similarity 73.4%;
Matches 1100; Conservative
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ORGANISM: Xenorhabdus
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RESULT 9 US-09-897-516-3855 ; Sequence 3855, Application US/09897516

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GENERAL INFORMATION:

APPLICANT: Corbin, David R.

APPLICANT: Goldman, Barry S.

APPLICANT: Hinkle, Gregory J.

APPLICANT: Huesing, Joseph E.

APPLICANT: Krasomil-Osterfeld, Karina C.

APPLICANT: Krasomil-Osterfeld, Karina C.

APPLICANT: Slater, Steven C.

APPLICANT: Slater, Steven C.

APPLICANT: Spiridonov, Sergei

TITLE OF INVENTION: Xenorihabdus sp. Genome Series Steven C.

CORRENT APPLICATION NUMBER: US/09/897,516

CURRENT FILING DATE: 2001-06-29

PRIOR APPLICATION NUMBER: US 60/215, 161

PRIOR FILING DATE: 2000-06-30
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SEQ ID NO 3855
LENGTH: 83080
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LOCATION: (7060)..(8658)
)9-897-516-3855
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RESULT 10
US-09-897-516-3856
Sequence 3856, Application US/09897516
GENERAL INFORMATION:
APPLICANT: Corbin, David R.
APPLICANT: Hinkle, Gregory J.
APPLICANT: Huesing, Joseph E.
APPLICANT: Krasomil-Osterfeld, Karina C.
APPLICANT: Malvar, Thomas M.
APPLICANT: Slater, Steven C.
APPLICANT: Slater, Steven C.
APPLICANT: Spiridonov, Sergei
FITTLE OF INVENTION: Xenorhabdus sp. Genome Sequences
FILE REFERENCE: 38-21(51847)B
CURRENT APPLICATION NUMBER: US/09/897,516

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Best Local Similarity
Matches 1100; Conser
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PRIOR APPLICATION NUMBER: US 60/
PRIOR FILING DATE: 2000-06-30
NUMBER OF SEQ ID NOS: 8409
SEQ ID NO 3856
LENGTH: 83080
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NAME/KEY: CDS
LOCATION: (8720)..(10027)
-09-897-516-3856
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ORGANISM: Xenorhabdus
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; LENGTH: 83080
; TYPE: DNA
; ORGANISM: Xenorhabdus sp.
; FEATURE:
; FEATURE: CDS
; LOCATION: (10097)..(11929)
US-09-897-516-3857
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                                                                                PRIOR APPLICATION NUMBER: US OPRIOR FILING DATE: 2000-06-30 NUMBER OF SEQ ID NOS: 8409 SEQ ID NO 3857 LENGTH: 83080
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GENERAL INFORMATION:
                                                                                                                                                    APPLICANT: Spiridonov, Sergei TITLE OF INVENTION: Xenorhabdus : FILE REFERENCE: 38-21(51847)B CURRENT APPLICATION NUMBER: US/09 CURRENT FILING DATE: 2001-06-29 CURRENT FILING DATE: 2001-06-29
                                                                                                                                                                                                                                                                                    APPLICANT: Corbin, David
APPLICANT: Goldman, Bar:
APPLICANT: Hinkle, Grego
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                                                                                                                                                                                                                          Goldman, Barry S.
Hinkle, Gregory J.
Huesing, Joseph E.
Krasomil-Osterfeld, I
Malvar, Thomas M.
Slater, Steven C.
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Pred. No. 1.9e-252;
0; Mismatches 356;
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APPLICANT: Corbin, David R.
APPLICANT: Goldman, Barry S.
APPLICANT: Hinkle, Gregory J.
APPLICANT: Huesing, Joseph E.
APPLICANT: Huesing, Joseph E.
APPLICANT: Krasomil-Osterfeld, Karina APPLICANT: Slater, Steven C.
APPLICANT: Spiridonov, Sergei
TITLE OF INVENTION: Xenorhabdus sp. Ge
FILE REFERENCE: 38-21(51847)B
CURRENT APPLICATION NUMBER: US/09/897,
CURRENT TILING DATE: 2001-06-29
PRIOR APPLICATION NUMBER: US 60/215, 1
PRIOR FILING DATE: 2000-06-30
NUMBER OF SEQ ID NOS: 8409
SEQ ID NO 3858
LENGTH: 83080
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LOCATION: (17551)..(19197)
-09-897-516-3858
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ORGANISM: Xenorhabdus
FEATURE:
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; TYPE: DNA; CORGANISM: Xenorhabdus sp.; FEATURE: ; FEATURE: ; NAME/KEY: CDS; LOCATION: (37881)..(38108) US-09-897-516-3859
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APPLICANT: Slater, Steven C.
APPLICANT: Spiridonov, Sergei
TITLE OF INVENTION: Xenorhabdus sp. Genome S
FILE REFERENCE: 38-21(51847)B
CURRENT APPLICATION NUMBER: US/09/897,516
CURRENT FILING DATE: 2001-06-29
PRIOR APPLICATION NUMBER: US 60/215, 161
PRIOR APPLICATION NUMBER: US 60/215, 161
PRIOR FILING DATE: 2000-06-30
NUMBER OF SEQ ID NOS: 8409
SEQ ID NO 3859
LENGTH: 83080
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US-09-897-516-3859
; Sequence 3859, Application US/09897516
; GENERAL INFORMATION:
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Best Local Similarity 73.4%;
Matches 1100; Conservative
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Huesing, Joseph E.
Krasomil-Osterfeld, Malvar, Thomas M.
Slater, Steven C.
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Pred. No. 1.9e-252;
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APPLICANT: COTBIN, David R.
APPLICANT: GOldman, Barry S.
APPLICANT: Hinkle, Gregory J.
APPLICANT: Huesing, Joseph E.
APPLICANT: Huesomil-Osterfeld, Karina
APPLICANT: Malvar, Thomas M.
APPLICANT: Slater, Steven C.
APPLICANT: Slater, Steven C.
APPLICANT: Spiridonov, Sergei
TITLE OF INVENTION: Xenorinabdus sp. Ge
FILE REFERENCE: 38-21(51847)B
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CURRENT FILING DATE: 2001-06-29
PRIOR APPLICATION NUMBER: US 60/215, 161
PRIOR FILING DATE: 2000-06-30
NUMBER OF SEO ID NOS: 8409
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ORGANISM: Xenorhabdus
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Pred. No. 1.9e-252;
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APPLICANT: Spiridonov, Sergei
TITLE OF INVENTION: Xenorhabdus sp. Genome S
FILE REFERENCE: 38-21(51847)B
CURRENT APPLICATION NUMBER: US/09/897,516
CURRENT FILING DATE: 2001-06-29
PRIOR APPLICATION NUMBER: US 60/215, 161
PRIOR FILING DATE: 2000-06-30
NUMBER OF SEQ ID NOS: 8409
SEQ ID NO 3861
LENGTH: 83080
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; NAME/KEY: CDS; LOCATION: (41695)..(42120)
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APPLICANT: Goldman, Barr
APPLICANT: Hinkle, Grego
APPLICANT: Huesing, Jose
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Hinkle, Gregory J.
Huesing, Joseph E.
Krasomil-Osterfeld,
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Search completed: December 15, 2001, 05:58:16 Job time: 12259 sec

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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Sequence 185, App	051	951	23,	e 23,	802,	ce 802,	1291,	equence 10329	ø	equence 18048	equence 1225,	equence 1	1452	equence 18351,	15063,	_	937, A	18350,	650,	19008,	19072,	22313,	7704	Sequence 148, App	352,	352,	289, A	11564	11774	17747.	49885	57118	908	17:	175385	7425 A	11944.	1. Appl	7. App	quence 2, Ap	Description		

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Sequence 7, Application US/09380826A
GENERAL INFORMATION:
APPLICANT: Chappel, Rod
TITLE OF INVENTION: LEPTOSPIRA PATHOGENS
FILE REFERENCE: DAVIET9.001APC
CURRENT APPLICATION NUMBER: US/09/380,826A
CURRENT FILING DATE: 1999-11-22
PRIOR APPLICATION NUMBER: PCT/AU98/00145
PRIOR PILING DATE: 1998-03-06
PRIOR FILING DATE: 1998-03-06
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                                                                                                 Query Match
Best Local S
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PRIOR FILING DATE: 1998-03-06
PRIOR APPLICATION NUMBER: AU PO5494/97
PRIOR FILING DATE: 1997-03-07
PRIOR FILING DATE: 1997-03-07
NUMBER OF SEQ ID NOS: 26
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 2
LENGTH: 22
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APPLICANT: Chappel, Rod
TITLE OF INVENTION: LEPTOSPIRA PATHOGENS
                                                                                                                                                                                                                                       SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 7 LENCTH: 22 TYPE: DNA
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Best Local Similarity
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CURRENT APPLICATION NUMBER: US/09/380,826A
CURRENT FILING DATE: 1999-11-22
                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: AU PO5494/97
PRIOR FILING DATE: 1997-03-07
NUMBER OF SEQ ID NOS: 26
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ORGANISM: Artificial Sequence
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GENERAL INFORMATION:
APPLICANT CORAPPEL, ROD
TITLE OF INVENTION: LEPTOSPIRA PATHOGENS
FILE REFERENCE: DAVIE79.001APC
CURRENT APPLICATION NUMBER: US/09/380,826A
CURRENT FILLING DATE: 1999-11-22
PRIOR APPLICATION NUMBER: PCT/AU98/00145
PRIOR TILING DATE: 1998-03-06
PRIOR FILING DATE: 1998-03-06
PRIOR APPLICATION NUMBER: AU PO5494/97
                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: ASPERGILLUS NIDULANS GENOME SEQUENCE AND USES THEREOF; FILE REFERENCE: 38-21(15485)C; CURRENT APPLICATION NUMBER: US/60/132,861; CURRENT FILING DATE: 1999-05-07; NUMBER OF SEQ ID NOS: 13800; SEQ ID NO 11944 LENCTH: 420; TYPE: DNA ORGANISM: Aspergillus nidulans US-60-132-861-11944
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               B
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                                                                                                                                                                   US-09-649-165A-7425/c : Sequence 7425, Application US/09649165A : GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 11944, Application US/60132861
GENERAL INFORMATION:
APPLICANT: Cao, Yongwei
APPLICANT: Timberlake, William E.
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Best Local Similarity
Matches 20; Conserv
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                          APPLICANT: Guegler, Karl J.
APPLICANT: Doyle, Martin
APPLICANT: Moniyama, Monika
APPLICANT: Wang, Xinhao
TITLE OF INVENTION: DETECTING EXPRESSION OF ARABIDOPSIS THALIANA GENES
FILE REFERENCE: PA-0019 US
CURRENT APPLICATION NUMBER: US/09/649,165A
CURRENT FILING DATE: 2000-08-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQ ID NOS: 26 SOFTWARE: FastSEQ for Windows Version 4.0
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ORGANISM: Leptospira
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                APPLICATION NUMBER:
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Pred. No. 1.1e+02;
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Pred. No.
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Best Local S
Matches 20
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SEQ ID NO 175385
LENGTH: 581
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APPLICANT: Kovalic,
APPLICANT: Liu, Jingdong
TITLE OF INVENTION:
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LENGTH: 530
                                                                                      SEQ ID NO 175385
LENGTH: 581
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Best Local Similarity
Query Match
                                           TYPE: DNA
ORGANISM: Arabidopsis thaliana
-09-684-016-175385
                                                                                                                                                         FILE OF INVENTION: Annotated Plant Genes FILE REFERENCE: 38-21(15097)D CURRENT APPLICATION NUMBER: US/09/684,016 CURRENT FILING DATE: 2000-10-10 PRIOR ADDITORATOR VICTORIO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: Annotated Plant Genes FILE REFERENCE: 38-21(15097)D CURRENT APPLICATION NUMBER: US/09/654,617 CURRENT FILING DATE: 2000-09-05
                                                                                                                                                                                               APPLICANT: Liu, Jin
TITLE OF INVENTION:
FILE REFERENCE:
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                                                                                                                       NUMBER OF SEQ ID NOS:
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OTHER INFORMATION: Incyte ID No:
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| 420 tgttggttcacaagatttgaca 441
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                                                                                                                                                                                                                              Jingdong
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Pred. No. 1
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Pred. No. 1
Score 18.8;
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FEATURE:

NAME/KEY: misc_feature

LOCATION: (1)..(1069)

OTHER INFORMATION: any n = a

NAME/KEY: misc_feature

LOCATION: (1)..(1069)

OTHER INFORMATION: 27496 : US-09-565-309A-57118
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NAME/KEY: misc_feature
; LOCATION: (1)..(649)
; OTHER INFORMATION: any n = a, g, c, t, unknown, or other
NAME/KEY: misc_feature
; LOCATION: (1)..(649)
; LOCATION: (1)..(649)
; OTHER INFORMATION: 27496:110925 (Clone Number:Unique Sequence Identifier)
US-09-565-309A-20908
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APPLICANT: ALEXANDROV, Nickolai; APPLICANT: BROVER, Vyacheslav
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; Sequence 20908, Application US/09565309A
; GENERAL INFORMATION:
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LENGTH: 649
TYPE: DNA
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SEQ ID NO 57118
LENGTH: 1069
TYPE: DNA
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Best Local Similarity
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              Query Match
Best Local Similarity
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APPLICANT: BROVER, Vyacheslav
TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTID
TITLE OF INVENTION: THEREBY
FILE REFERENCE: 2750-0853P
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TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS AND CORRESPONDING POLYPEPTID
TITLE OF INVENTION: THEREBY
FILE REFERENCE: 2750-0853P
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CURRENT FILING DATE: 2000-05-05
NUMBER OF SEQ ID NOS: 68449
                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/09/565,309A CURRENT FILING DATE: 2000-05-05
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                85.5%;
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Score 18.8; D
Pred. No. 1.2e
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Pred. No. 1.1e+02;
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                                 Length 1069;
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NAME/KEY: misc_feature
LOCATION: (1):.(1072)
OTHER IMFORMATION: any n = a
NAME/KEY: misc_feature
LOCATION: (1):.(1072)
OTHER IMFORMATION: 27496 : 5'
US-09-565-309A-49885
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US-09-565-309A-49885
US-09-684-016-117747; App; Sequence 117747, App; GENERAL INFORMATION:
                                                                                                В
                                                                                                                                                                                                                       ; TYPE: DNA; ORANISM: Arabidopsis thaliana; OTHER INFORMATION: unsure at all n locations US-09-654-617-117747
                                                                                                                                                                                                                                                                         Annotated Plant Genes
CURRENT APPLICATION NUMBER: US/09/654,617
CURRENT FILING DATE: 2000-09-05
MUMBER OF SEQ ID NO 117747
LENGTH: 1170
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Best Local Similarity
Matches 20; Conser
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SEQ ID NO 49885
LENGTH: 1072
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Best Local S
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APPLICANT: BROVER, VYACHOSLAV
TITLE OF INVENTION: SEQUENCE-DETERMINED DNA FRAGMENTS
TITLE OF INVENTION: THEREBY
FILE REFERENCE: 2750-0853P
CURRENT APPLICATION NUMBER: US/09/565,309A
CURRENT FILING DATE: 2000-05-05
                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Kovalic, APPLICANT: Liu, Jingdong
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                           Application US/09684016
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90.9%;
  David
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Pred. No. 1.2e
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Pred. No. 1.2e+02;
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Annotated Plant Genes
CURRENT APPLICATION NUMBER: US/097)D
CURRENT FILING DATE: 2000-10-10
PRIOR APPLICATION NUMBER: 09/654,617
PRIOR FILING DATE: 2000-09-05
PRIOR FILING DATE: 2000-09-05
COURTENT F
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; LOCATION: (1)..(1128)
; OTHER INFORMATION: unsure at all n
US-09-684-016-117747
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; ORGANISM: Aspergillus nidulans
US-60-138-103-11564
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GENERAL INFORMATION:
APPLICANT: Cao, Yongwei
APPLICANT: Timberlake, William E.
TITLE OF INVENTION: ASPERGILUS NIDULANS GENOME SEQUENCE AND USES THEREOF
FILE REFERENCE: 38-10(15485)
CURRENT APPLICATION NUMBER: US/60/138,103
CURRENT FILING DATE: 1999-06-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS: 28006
SEQ ID NO 11564
LENGTH: 1429
NUMBER OF SEQ ID NOS:
SEQ ID NO 5289
LENGTH: 5874
TYPE: DNA
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Best Local Similarity
Matches 20; Conserv
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Best Local
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                                                                                                                                          APPLICANT: Cao, Yongwei
APPLICANT: Ghodssi, Azita
APPLICANT: Hinkle, Gregory J.
APPLICANT: McIninch, James
APPLICANT: Timberlake, William E.
APPLICANT: Yu, Jeebyuk
TITLE OF INVENTION: Emericella nidulans Genome Sequence
FILE REFERENCE: 38-10(15498)A
CURRENT APPLICATION NUMBER: US/09/404,520
CURRENT FILING DATE: 1999-09-23
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Pred. No. 1.2e+02;
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Pred. No. 1.2e+02;
0; Mismatches 2;
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APPLICANT: Last, Robert L.
APPLICANT: Levin, Irena M.
APPLICANT: Norris, Susan R.
APPLICANT: Norris, Susan R.
APPLICANT: Parnell, Laurence D.
APPLICANT: Wiegand, Roger C.
APPLICANT: Wiegand, Roger C.
TITLE OF INVENTION: PLANT POLYMORPHIC N
FILE REFERENCE: 38-10(15493)B
CURRENT APPLICATION NUMBER: US/09/534,859
CURRENT FILING DATE: 2000-03-29
NUMBER OF SEQ ID NOS: 1127
SEQ ID NO 352
LENGTH: 92624
TYPE: DNA
ORGANISM: Arabidopsis thaliana
US-09-534-859-352
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US-09-534-859-352
Search completed: December 15, 2001, 05:42:55 Job time: 15820 sec
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GENERAL INFORMATION:
APPLICANT: Bush, David F.
                                                                                                                                                                                       Ouery Match 85.5%; Score 18.8; DB 20; Length 92624; Best Local Similarity 90.9%; Pred. No. 2e+02; Matches 20; Conservative 0; Mismatches 2; Indels 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ouery Match 85.5%; Score 18.8; DB 18; Length 5874; Best Local Similarity 90.9%; Pred. No. 1.4e+02; Matches 20; Conservative 0; Mismatches 2; Indels 0;
                                                                                                     PLANT POLYMORPHIC MARKERS AND USES THEREOF
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Perfect score:
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1: /cgn2_6/ptodata/1/pna/U
2: /cgn2_6/ptodata/1/pna/U
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4: /cgn2_6/ptodata/1/pna/U
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/cgn2_6/ptodata/1/pna/US06_NEW_COMB.seq:*
/cgn2_6/ptodata/1/pna/US07_NEW_COMB.seq:*
/cgn2_6/ptodata/1/pna/US08_NEW_COMB.seq:*
/cgn2_6/ptodata/1/pna/US09_NEW_COMB.seq:*
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US-09-985-402-1036
US-09-815-264-26466
US-09-815-264-26466
US-09-815-264-23140
US-09-815-264-3319
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Compugen Ltd
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                                                                    Sequence 4668, Ap
Sequence 210, App
Sequence 211824,
Sequence 2773, Ap
Sequence 1036, A
Sequence 27340, A
Sequence 26466, A
Sequence 63319, A
Sequence 33140, A
Sequence 32140, A
Sequence 53250, A
Sequence 76653, A
Sequence 76653, A
Sequence 71043, A
Sequence 71034, A
Sequence 71944, A
Sequence 719466, A
Sequence 719466, A
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18469,
238, AI
2094, 1
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   NUMBER OF
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	15.8	16	16	16	16	16.2	16.2	16.2	16.2	16.2	16.2	16.2	16.2	16.2	16.2	16.2	16.2	16.2	16.2	16.2
	71.8	72.7	72.7	72.7	72.7	73.6	73.6	73.6	73.6	73.6	73.6	73.6	73.6	73.6	73.6	73.6	73.6	73.6	73.6	73.6
	270	39365	37658	1381	610	36227	32005	26601	10839	9723	7856	6237	3603	1501	1501	1501	1457	1251	845	461
•	6	6	6	6	6	σ	6	σ	6	6	δ	6	φ	σ	δ	σ	σ	σ	6	σ
ALIGNMENTS	US-09-985-678-42368	US-09-815-264-79204	US-09-815-264-74259	US-09-815-264-108378	US-09-985-678-213635	US-09-815-264-68619	US-09-815-264-73015	US-09-815-264-74669	US-09-815-264-78946	US-09-815-264-75693	US-09-815-264-86401	US-09-815-264-72243	US-09-815-264-80803	US-09-815-264-43531	US-09-815-264-39776	US-09-815-264-39775	US-09-815-264-60838	US-09-898-888A-9009	US-09-922-279A-1268	US-09-849-526A-12135
	Sequence 42368,	Sequence 79204,	Sequence 74259,	Sequence 108378	Sequence 213635	Sequence 68619,	Sequence 73015,	Sequence 74669,	Sequence 78946,	Sequence 75693,	Sequence 86401,	Sequence 72243,	Sequence 80803,	Sequence 43531,	Sequence 39776,	Sequence 39775,	Sequence 60838,	Sequence 9009,	Sequence 1268,	Sequence 12135,
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US-09-620-394B-4668
Sequence 4668, Application US/09620394B
GENERAL INFORMATION:
APPLICANT: ALEXANDROV, Nickolai
APPLICANT: ALEXANDROV, Nickolai
APPLICANT: BROVER, Vyacheslav
TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptid
TITLE OF INVENTION: Thereby
FILE REFERENCE: 2750-1067P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; OTHER INFORMATION: any n = a, g, c, t, u, NAME/KBY: misc_feature; LOCATION: 1..1069; OTHER INFORMATION: Ceres Seq. ID 1393173 US-09-620-394B-4668
                                                                                                                                                              Sequence 520, Application US/09924035A GENERAL INFORMATION: APPLICANT: Grlach, Jrn
                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
Matches 20; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/620,394B
CURRENT FILING DATE: 2000-07-21
NUMBER OF SEQ ID NOS: 9131
SEQ ID NO 4668
              TITLE OF INVENTION: Expressed Sequences of TITLE OF INVENTION: thallana FILE REFERENCE: 2011US CURRENT APPLICATION NUMBER: US/09/924,035A CURRENT FILING DATE: 2000-08-11 PRIOR APPLICATION NUMBER: US 60/148,784 PRIOR FILING DATE: 1999-08-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA
ORGANISM: Arabidopsis thaliana
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: misc_feature
LOCATION: 1..1069
                                                                                                                                                                                                                                                                                                      690 tgttggttcacaagatttgaca
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SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                             90.9%;
                                                                                                                                        Sequences of Arabidopsis
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Pred. No. 1
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APPLICANT: Liu, Jingdong
TITLE OF INVENTION: Annotated Plant Genes
FILE REFERENCE: 16517.255/38-21(15097)F
CURRENT APPLICATION NUMBER: US/09/985,678
CURRENT FILING DATE: 2001-11-05
PRIOR APPLICATION NUMBER: US 09/304,517
PRIOR APPLICATION NUMBER: US 09/304,517
PRIOR FILING DATE: 1999-05-06
NUMBER OF SEQ ID NOS: 295529
SEQ ID NO 211824
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 3
US-09-985-678-211824
Sequence 211824, Application US/09985678
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Cheikh, Nordine
Liu, Jingdong
APPLICANT: Liu, Jingdong
APPLICANT: Cheikh, Annotated Plant Gen
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NAME/KEY: misc_feature

; LOCATION: (1)...(249):

OTHER INFORMATION: n = A,T,C or

US-09-924-035A-520
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LOCATION: (1)..(622)
COTHER INFORMATION: unsure at all n locations
US-09-985-678-211824
                            APPLICANT: L1u, Chenghua
APPLICANT: Drmanac, Radoje T.
TITLE OF INVENTION: NOVel Contigs Obtained
TITLE OF INVENTION: From Various Libraries
FILE REFERENCE: 785
CURRENT APPLICATION NUMBER: US/09/922,279A
CURRENT FILING DATE: 2001-08-03
PRIOR APPLICATION NUMBER: 09/491,404
PRIOR FILING DATE: 2000-01-25
NUMBER OF SEO ID NOS: 3796
                                                                                                                                                                                                                                                                                     US-09-922-279A-2773; Sequence 2773, Application US/09922279A; GENERAL INFORMATION:
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SEQ ID NO 520
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Best Local Similarity
Matches 19; Conserva
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Best Local Similarity
Matches 19; Conserv
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TYPE: DNA
ORGANISM: Arabidopsis thaliana
                                                                                                                                                                                      APPLICANT: Tang, Yuanhua T.
APPLICANT: Tillinghast, Jo
APPLICANT: Sinku, Ankura
APPLICANT: Liu, Chenghua
APPLICANT: Drmanac, Radoje
SOFTWARE: pt_SP_genes Version 1.0 EQ ID NO 2773
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Glycine max
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                    520 tgatggatctcaagctttgata 541
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86.4%;
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86.4%;
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Pred. No. 81;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
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pred. No. 87;
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NAME/KEY: sig_peptide
LOCATION: (1991)...(956)
THER INFORMATION: this location contains the signal peptide sequence,
OTHER INFORMATION: MLYLLLFPGVSYLRSLFLGRPIGPGITSDFTLILFSNLLDSWPLS, Run with Signal
NAME/KEY: misc_feature
LOCATION: (605)...(1902)
OTHER INFORMATION: similar to gi5931821 in the genepept database release 114,
OTHER INFORMATION: Run with FASTXY 3.3t00, default parameters
US-09-922-279A-2773
APPLICANT: [list inventors here]
TITLE OF INVENTION: [list title here]
FILE REFERENCE: [list docket ID here]
CURRENT APPLICATION NUMBER: US/09/982,402
CURRENT FILING DATE: 2001-10-17
NUMBER OF SEQ ID NOS: 3553
SOFTWARE: PERL PROGRAM
SEQ ID NO 1036
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US-09-815-264-82805
                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; ORGANISM: Oryza sativa
US-09-815-264-82805
                                                                                                                                                                                                       US-09-982-402-1036
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: Genomic Plant Sequences And Uses Thereof FILE REFERENCE: 38-21(51237)G CURRENT APPLICATION NUMBER: US/09/815,264 CURRENT FILING DATE: 2001-03-23 PRIOR APPLICATION NUMBER: US 09/620,392 PRIOR APPLICATION NUMBER: US 09/620,392 PRIOR APPLICATION NUMBER: US 09/702,134 PRIOR FILING DATE: 2000-10-31 NUMBER OF SEQ ID NO 82805 SEQ ID NO 82805 LENGTH: 46891
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 82805, Application US/09815264
GENERAL INFORMATION:
APPLICANT: Boukharov, Andrey A.
APPLICANT: Cao, Yongwei
APPLICANT: Dotson, Stanton B.
APPLICANT: Koshi, Jeffrey M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches 19;
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Best Local Similarity
                                                                                                                                                           Sequence 1036, Application US/09982402 GENERAL INFORMATION:
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Best Local Similarity
Matches 19; Conserv
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APPLICANT:
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                                                                                                                                                                                                                                                                                45218 tgttgcaacacaagacttgata 45239
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
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Koshi, Jeffrey M.
Kovalic, David K.
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86.4%;
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Pred. No. 96;
0; Mismatches
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Pred. No. 1.3e
0; Mismatches
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1.3e+02;
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ORGANISM: Zea mays

FEATURE:

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APPLICANN: RUIF, Thomas G.
APPLICANN: RUIF, Thomas G.
APPLICANT: Shukla, Hridayabhiranjan
APPLICANT: Shukla, Hridayabhiranjan
APPLICANT: Shukla, Hridayabhiranjan
TITLE OF INVENTION: Nucleic Acid Molecules And
TITLE OF INVENTION: Nucleic Sci (51930)B]
CURRENT APPLICATION NUMBER: US/09/849,526A
CURRENT FILING DATE: 2001-05-07
PRIOR APPLICATION NUMBER: US/09/202,214
PRIOR APPLICATION NUMBER: US/09/64,617
PRIOR APPLICATION NUMBER: US/09/64,617
PRIOR APPLICATION NUMBER: US/09/64,016
PRIOR APPLICATION NUMBER: US/09/684,016
PRIOR APPLICATION NUMBER: US/09/684,016
PRIOR APPLICATION NUMBER: US/09/684,016
PRIOR APPLICATION NUMBER: US/09/684,016
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; OTHER INFORMATION: Incyte ID No: 700196524H1
US-09-982-402-1036
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SEQ ID NO 27340
LENGTH: 551
                                                                                                                                                              GENERAL INFORMATION:
APPLICANT: Boukharov, Andrey A.
APPLICANT: Cao, Yongwei
APPLICANT: Dotson, Stanton B.
APPLICANT: Koshi, Jeffrey M.
APPLICANT: Kovalic, David K.
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Best Local
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Best Local :
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                    TITLE OF INVENTION: Genomic Plant Sequences FILE REFERENCE: 38-21(51237)G CURRENT APPLICATION NUMBER: US/09/815,264 CURRENT FILING DATE: 2001-03-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: US 09/816,660 PRIOR FILING DATE: 2001-03-26
                                                                                                    APPLICANT:
                                                                                                                                               APPLICANT:
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nes 18; Conserv
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APPLICATION NUMBER: US 09/620,392
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Wu, Wei
                                                                                                                                               Liu, Jingdong
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Pred. No. 1
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                                                                                And Uses Thereof
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 551;
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; ORGANISM: Oryza sativa
US-09-815-264-63319
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US-09-815-264-26466
                                                                                                                                                                                                                                                      US-09-815-264-91672
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-09-815-264-63319
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APPLICANT: Cao, Yongwei
APPLICANT: Dotson, Stanton B.
APPLICANT: Koshi, Jeffrey M.
APPLICANT: Kovalic, David K.
APPLICANT: Liu, Jingdong
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Best Local Similarity 90.0
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SEQ ID NO 26466
LENGTH: 849
TYPE: DNA
                                                                                                                                                                                                                  Sequence 91672, Application US/09815264 GENERAL INFORMATION:
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SEQ ID NO 63319
LENGTH: 1046
TYPE: DNA
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Best Local Similarity 90.0%;
Matches 18; Conservative
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APPLICANT:
APPLICANT:
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PRIOR APPLICATION NUMBER: US
PRIOR FILING DATE: 2000-10-31
APPLICANT: Wu, Wei
ITITE OF INVENTION: Genomic Plant Sequences and Uses Thereof
FILE REFERENCE: 38-21(51237)G
CURRENT APPLICATION NUMBER: US/09/815,264
CURRENT FILING DATE: 2001-03-23
                                                                                                                                         APPLICANT: Boukharov, Andrey A. APPLICANT: Cao, Yongwei APPLICANT: Dotson, Stanton B. APPLICANT: Koshi, Jeffrey M.
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Koshi, Jeffrey M.
Kovalic, David K.
                                                                                                         Liu, Jingdong
                                                                                    McIninch, James
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90.08;
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Pred. No. 1.4e+02;
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US-09-815-264-91672
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; ORGANISM: Oryza sativa
US-09-815-264-23140
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SEQ ID NO 91672
LENGTH: 1246
TYPE: DNA
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Best Local Similarity
Matches 18; Conserv
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PRIOR FILING DATE: 2000-10-31
NUMBER OF SEQ ID NOS: 109669
SEQ ID NO 23140
LENGTH: 1501
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GENERAL INFORMATION:
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PRIOR FILING DATE: 2000-07-19
PRIOR APPLICATION NUMBER: US 09/702,134
PRIOR FILING DATE: 2000-10-31
                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
Matches 18; Conserv
                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                           Sequence 46882, Application US/09815264
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                                                                                             APPLICANT:
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APPLICANT: Wu, Wei
TITLE OF INVENTION: Genomic Plant Sequences FILE REFERENCE: 38-21(51237)G
CURRENT APPLICATION NUMBER: US/09/815,264
                                                                  APPLICANT:
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Koshi, Jeffrey M.
Kovalic, David K.
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                                                                                           Dotson, Stanton B.
Koshi, Jeffrey M.
Kovalic, David K.
                                                             McIninch, James
                                                                                 Liu, Jingdong
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90.0%;
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Pred. No. 1
                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                          Score 16.8; DB 6; Pred. No. 1.4e+02;
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1.4e+02;
ches 2;
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                                       And
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: NAME/KEY: unsure
: LOCATION: (1)..(1501)
: OTHER INFORMATION: unsure at all n locations
US-09-815-264-46882
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PRIOR FILING DATE: 2000-07-19
PRIOR APPLICATION NUMBER: US 09/702,134
PRIOR FILING DATE: 2000-10-31
NUMBER OF SEQ ID NOS: 109669
SEQ ID NO 46882
LENGTH: 1501
                                                                                                                                                                                                                                                                                                                                                     ; ORGANISM: Oryza sativa
US-09-815-264-53250
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                                                                                                                                RESULT 14
US-09-815-264-76889
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Best Local
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SEQ ID NO 53250
LENGTH: 1501
TYPE: DNA
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                                                                                             Sequence 76889, Application US/09815264 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILE REFERENCE: 38-21(51237)G
CURRENT APPLICATION NUMBER: US/09/815,264
CURRENT FILING DATE: 2001-03-23
PRIOR APPLICATION NUMBER: US 09/620,392
PRIOR FILING DATE: 2000-07-19
PRIOR FILING DATE: 1000-07-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Liu, Jingdong
APPLICANT: McIninch, James
APPLICANT: Wu, Wei
TITLE OF INVENTION: Genomic Plant Sequences And Uses Thereof
FILE REFERENCE: 38-21(51237)9
                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: US 09/702,134 PRIOR FILING DATE: 2000-10-31
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                           APPLICANT: Boukharov, Andrey A.
APPLICANT: Cao, Yongwei
APPLICANT: Dotson, Stanton B.
APPLICANT: Koshi, Jeffrey M.
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Koshi, Jeffrey M.
Kovalic, David K.
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              Dotson, Stanton B
Koshi, Jeffrey M.
Kovalic, David K.
Liu, Jingdong
                                                                                                                                                                                                                                                                       Conservative
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90.0%;
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Pred. No. 1.
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                                                                                                                                                                                                                                                                                           Score 16.8;
Pred. No. 1.
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1.4e+02;
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APPLICANT:

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PRIOR APPLICATION NUMBER: US 09/620,392
PRIOR FILING DATE: 2000-07-19
PRIOR APPLICATION UNMBER: US 09/702,134
PRIOR FILING DATE: 2000-10-31
NUMBER OF SEQ ID NOS: 109669
SEQ ID NO 81045
LENGTH: 12081
TYPE: DNA
ORGANISM: OTYZA SALÍVA
US-09-815-264-81045
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: McIninch, James

APPLICANT: Wu, Wei

TITLE OF INVENTION: Genomic Plant Sequences And Uses Thereof

FILE REFERENCE: 38-21(51237)G

CURRENT APPLICATION NUMBER: US/09/815,264

CURRENT FILING DATE: 2001-03-23

PRIOR APPLICATION NUMBER: US 09/620,392

PRIOR APPLICATION NUMBER: US 09/620,392

PRIOR FILING DATE: 2000-07-19

PRIOR APPLICATION NUMBER: US 09/702,134

PRIOR FILING DATE: 2000-10-31

PRIOR FILING DATE: 2001-10-31

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NAME/KEY: unsure

; LOCATION: (1). (6940)

; OTHER INFORMATION: unsure at all n locations

US-09-815-264-76889
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Search completed: December 15, 2001, 05:58:19 Job time: 12262 sec
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Best Local Similarity 90.0
Matches 18; Conservative
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Best Local Similarity 90.0
Matches 18; Conservative
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TITLE OF INVENTION: Genomic Plant Sequences And Uses Thereof
FILE REFERENCE: 38-21(51237)G
CURRENT APPLICATION NUMBER: US/09/815,264
CURRENT FILING DATE: 2001-03-23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
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TYPE: DNA
ORGANISM: Oryza sativa
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3165 ttggaccaaaagatttgata 3184
                                                                                                               Dotson, Štanton B.
Koshi, Jeffrey M.
Kovalic, David K.
Liu, Jingdong
McIninch, James
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cao, Yongwei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Application US/09815264
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      76.4%; Score 16.8; DB 6; 90.0%; Pred. No. 1.7e+02; 91:00 0: Mismatches 2;
                                                                                                                                                                                                                                    76.4%; Score 16.8; DB 6; 90.0%; Pred. No. 1.7e+02;
                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                Length 12081;
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Minimum DB seq length: 0
Maximum DB seq length: 2000000000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Title:
Perfect score:
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Maximum Match 1008
Listing first 45 summaries
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Copyright (c) 1993 - 2000 Compugen Ltd
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          tgttgga 7
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15.407 Million cell updates/sec
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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100.0 100.0

Sequence Sequence

Sequence

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US-09-065-775-44
; Sequence 44, Application US/09065775A
; GENERAL INFORMATION:
; APPLICANT: Cen, Hui
; APPLICANT: Sun, Shaojian
; TITLE OF INVENTION: Function-Based Gene Discovery
; FILE REFERENCE: 59818
; CURRENT APPLICATION NUMBER: US/09/065,775A
; CURRENT FILING DATE: 1998-04-24
; NUMBER OF SEQ ID NOS: 105
; SOFTMARE: Patentin Ver. 2.1
; SEQ ID NO 44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Chappel, Rod
TITLE OF INVENTION: LEPTOSPIRA PATHOGENS
FILE REFERENCE: DAVIET9.001APC
CURRENT APPLICATION NUMBER: US/99/380,826A,
CURRENT FILING DATE: 1999-11-22
PRIOR APPLICATION NUMBER: PCT/AU98/00145
PRIOR FILING DATE: 1998-03-06
PRIOR APPLICATION NUMBER: AU PO5494/97
PRIOR APPLICATION NUMBER: AU PO5494/97
PRIOR FILING DATE: 1997-03-07
NUMBER OF SEQ ID NOS: 26
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 4
LENGTH: 7
                                             Query Match
Best Local Similarity
Thes 7; Conserv
                                                                                                                                       ; OTHER INFORMATION: Description of Artificial Sequence: synthetic; OTHER INFORMATION: oligonucleotide primers US-09-065-775-44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
"heres 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; FEATURE:
; OTHER INFORMATION: Primer for L. fainei
US-09-380-826A-4
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                                                                                                                                                                                           TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Artificial Sequence
                                                                                                                                                                                      FEATURE:
                                                                                                                                                                                                                                   LENGTH:
                1 tgttgga 7
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tgttgga 9
                                                             Conservative
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                                                                           100.0%; Score 7; 100.0%; Pred. No.
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US-09-341-700A-421
US-09-341-700A-1193
US-09-504-231A-1407
                                                              0;
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Pred. No. 2.5
0; Mismatches
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                                                              Mismatches
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2.5e+09;
                                                                             DB 14; 1
1.9e+09;
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421, App
1193, Ap
1407, Ap
                                                              Gaps
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; SOFTWARE: PatentIn Ver.
; SEQ ID NO 149
; LENGTH: 10
; TYPE: DNA
; ORGANISM: Homo sapiens
PCT-US00-24784-149
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Best Local Similarity
Tatches 4; Conserve
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.US-09-916-466-170
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; Sequence 149, Application PC/TUS0024784
; GENERAL INFORMATION:
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Best Local Similarity
Thehes 7; Conserv
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                                                                           Sequence 47, Application PC/TUS0103620 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Chew, Anne
APPLICANT: Choi, Julie Y.
APPLICANT: Choi, Julie Y.
TITLE OF INVENTION: DRUG TARGET ISOGENES:
TITLE OF INVENTION: POLYMORPHISMS IN THE UNCOUPLING PROTEIN 3
TITLE OF INVENTION: (MITOCHONDRIAL, PROTON CARRIER) GENE
FILE REFERENCE: UCP3-0009PCT
CURRENT APPLICATION NUMBER: PCT/US00/24784
CURRENT FILING DATE: 2000-09-08
PRIOR APPLICATION NUMBER: MRH009
PRIOR FILING DATE: 1999-09-08
ROTELAPET DEFETTO VET 21
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APPLICANT:
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APPLICANT: Akhtar, Saghir
APPLICANT: Akhtar, Saghir
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or conditions Rel
TITLE OF INVENTION: Levels of Epidermal Growth Factor Receptors
FILE REFERENCE: MBHB00-958-J (400/032)
CURRENT APPLICATION NUMBER: US/09/916,466
CURRENT FILING DATE: 2001-07-25
NUMBER OF SEQ ID NOS: 446
APPLICANT: Genaissance Pharmaceuticals, Inc. APPLICANT: Chew, Anne APPLICANT: Koshy, Beena APPLICANT: Koshy, Beena APPLICANT: Choi, Julie Y.
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Denton, R. Rex
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Pred. No. 1.9e+09;
3; Mismatches 0;
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Pred. No.
                                                                                                                                                                                                                                                                            Mismatches
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APPLICANI: LUCULI, ....,
APPLICANI: Kazemi, Amir
APPLICANT: Koshy, Beena
FITLE OF INVENTION: Haplotypes of the BDKRB2 Gene
FILE REFERENCE: MMH-0740PCT BDKRB2
CURRENT APPLICATION NUMBER: PCT/US01/18811A
CURRENT FILING DATE: 2001-06-11
PRIOR APPLICATION NUMBER: 60/210,575
PRIOR APPLICATION NUMBER: 60/210,575
PRIOR APPLICATION NUMBER: 50/210,575
PRIOR APPLICATION NUMBER: 50/210,575
PRIOR APPLICATION NUMBER: 50/210,575
LENGTH: 10 DATE: 2000-06-09
SOFTWARE: PATENTIN VET. 2.1
SEQ ID NO 45
LENGTH: 10
TYPE: DNA
ORGANISM: Homo sapiens
PCT-US01-18811A-45
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; TYPE: DNA
; ORGANISM: Homo sapiens
PCT-US01-03620-47
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Best Local Similarity 100.
Matches 7; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
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APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Genaissance Pharmaceuticals, Inc. APPLICANT: Bentivegna, Steven C. APPLICANT: Bieglecki, Karyn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILE REFERENCE: MWH-0045 AGTR2
CURRENT APPLICATION NUMBER: PCT/US01/03620
CURRENT FILING DATE: 2001-02-02
PRIOR APPLICATION NUMBER: 60/179,922
PRIOR FILING DATE: 2000-02-03
NUMBER OF SEQ ID NOS: 74
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TITLE OF INVENTION: DRUG TARGET ISOGENES: POLYMORPHISMS IN THE ANGIOTENSIN
TITLE OF INVENTION: RECEPTOR 2 GENE
         CORRESPONDENCE ADDRESS:
ADDRESSEE: Vinson & Elkins
STREET: 1455 Pennsylvania Avenue, N.W.
                                                                                 APPLICANT: Beattie, Kenneth L.

TITLE OF INVENTION: Microfabricated, Flowthrough Porous
TITLE OF INVENTION: Apparatus for Discrete Detection of Binding Reactions
NUMBER OF SEQUENCES: 17
                                                                                                                                                                                                                                                                                                                                                                                         y Match 100.0%; Score 7; DB 1; Local Similarity 100.0%; Pred. No. 4.9e+hes 7; Conservative 0; Mismatches
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Washington
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Bieglecki, Karyn
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100.0%; Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: Genzyme Corporation
APPLICANT: Roberts, Bruce L.
APPLICANT: Shankara, Srinivas
TITLE OF INVENTION: PREPARATION AND USE OF SUPERIOR VACCINES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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EARLIER
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APPLICATION NUMBER: US 08/141
FILING DATE: 28-OCT-1994
ATTORNEY/AGENT INFORMATION:
NAME: Sanzo, Michael A.
REGISTRATION NUMBER: 36,912
REFERENCE/DOCKET NUMBER: "HARC'
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)639-65400
                                                                                                                                                                                               EARLIER FILING DATE: 1998-06-19 EARLIER APPLICATION NUMBER: 60/ EARLIER FILING DATE: 1998-06-19
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EARLIER APPLICATION NUMBER: 60/090,041
                                                                                                                                                                                                                                                                                                                                                                                            EARLIER APPLICATION NUMBER: 60/090,040
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                             EARLIER APPLICATION NUMBER: 60/089,993 EARLIER FILING DATE: 1998-06-19
                                                                                                                                                        EARLIER APPLICATION NUMBER: 60/090,035 EARLIER FILING DATE: 1998-06-19
                                                                                                                                                                                                                                                                                   EARLIER APPLICATION NUMBER: 60/089,853
EARLIER FILING DATE: 1998-06-19
                                                                                                                                                                                                                                                                                                                                 EARLIER FILING DATE: 1998-06-19
                                                                                                                                                                                                                                                                                                                                                                                                                     EARLIER APPLICATION NUMBER: 60/090,039 EARLIER FILING DATE: 1998-06-19
                                                                                                                                                                                                                                                                   EARLIER APPLICATION NUMBER: 60/089,997
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                           EARLIER
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COMPUTER READABLE FORM:

COMPUTER FLOPPY disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: .PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.25
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APPLICATION NUMBER: PC
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STRANDEDNESS: both
TOPOLOGY: linear
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                                                               APPLICATION NUMBER: 60/089,992 FILING DATE: 1998-06-19
                                           APPLICATION NUMBER: 60/090,072
APPLICATION NUMBER: 60/089,878
                      FILING DATE: 1998-06-19
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100.0%; Pred. No.
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thes 0;
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EARLIER FILING DATE: 1998-06-19
EARLIER APPLICATION NUMBER: 60/090,000

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; ORGANISM: Homo sapiens
PCT-US99-13800-679
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EARLIER FILING DATE: 1998-12-08
NUMBER OF SEQ ID NOS: 2138
SOFTWARE: FRASTSEQ for Windows Version 3.0
SEQ ID NO 679
                                                                                                                                                                                                                                                             Sequence 13, Application US/08141969
GENERAL INFORMATION:
APPLICANT: Beattie, Kenneth L.
TITLE OF INVENTION: Microfabricated, Flowthrough Porous
TITLE OF INVENTION: Apparatus for Discrete Detection of
NUMBER OF SEQUENCES: 17
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Best Local Similarity
Matches 7; Conserv
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EARLIER APPLICATION NUMBER: 60/
EARLIER FILING DATE: 1998-06-19
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EARLIER FILING DATE: 1998-06-19
EARLIER APPLICATION NUMBER: 60/089,994
EARLIER FILING DATE: 1998-06-19
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DDS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 tgttgga 7
|||||||
3 tgttgga 9
                                                                                                                                                               STREET: 1455 Peni
CITY: Washington
                                                                                                            COUNTRY: U.S.A. ZIP: 20004-1008
                                                                                                                                                      STATE: D.C.
                                                                                                                                                                                          ADDRESSEE: Vinson & Elkins
STREET: 1455 Pennsylvania Avenue, N.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ER APPLICATION NUMBER: 60/090,044
ER FILING DATE: 1998-06-19
ER APPLICATION NUMBER: 60/089,844
ER FILING DATE: 1998-06-19
ER APPLICATION NUMBER: 60/090,080
ER FILING DATE: 1998-06-19
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APPLICATION NUMBER: 60/090,042
FILING DATE: 1998-06-19
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FILING DATE: 1998-06-19
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                                                                                                                                                                                                                                                                                       Binding Reactions
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US-09-213-932A-13/c
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Best Local Similarity
Matches 7; Conser
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INFORMATION FOR SEQ ID NO:
Query Match
Best Local Similarity
                                                                                                                                                                      INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)639-6500
TELEFAX: (202)639-6604
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/08 FILING DATE: 28-OCT-1993 CLASSIFICATION: 435 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS: LENGTH: 10 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION:
                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 10 base pairs
TYPE: nucleic acid
STRANDEDNESS: both
                                                                                                                                                                                                                                      FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Sanzo, Michael A.
REGISTRATION NUMBER: 36,912
REFERENCE/DOCKET NUMBER: HAP
                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA:
                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)639-6500
                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
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STRANDEDNESS: both
TOPOLOGY: linear
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REGISTRATION NUMBER: 36,912
REFERENCE/DOCKET NUMBER: HAI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: Washington
                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET:
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1455 Pennsylvania Avenue, N.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               U.S.A.
                                                                                                                                                                                        (202)639-6604
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   100.0%;
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Pred. No.
     Score 7;
Pred. No.
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     DB 16;
4.9e+05;
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CURRENT APPLICATION NUMBER: US/09/335,032
CURRENT FILING DATE: 1999-06-16
PRIOR APPLICATION NUMBER: US 60/035,917
PRIOR FILING DATE: 1997-01-23
PRIOR FILING DATE: 1997-01-23
PRIOR APPLICATION NUMBER: US 09/012,031
PRIOR FILING DATE: 1998-01-22
NUMBER OF SEQ ID NOS: 12219
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 8949
LENGTH: 10
TYPE: DNA
ORGANISM: Saccharomyces cerevisiae
US-09-335-032-8949
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Best Local Similarity
Thes 7; Conserve
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APPLICANT: Vogelstein, Bert
APPLICANT: Vogelstein, Bert
APPLICANT: Kinzler, Kenneth
TITLE OF INVENTION: Characterization of the Yeast
TITLE OF INVENTION: Transcriptome
FILE REFERENCE: 01107.78572
CURRENT APPLICATION NUMBER: US/09/335,032
CURRENT FILING DATE: 1999-06-16
PRIOR APPLICATION NUMBER: US 60/035,917
PRIOR APPLICATION NUMBER: US 60/035,917
PRIOR APPLICATION NUMBER: US 60/035,917
PRIOR FILING DATE: 1997-01-23
PRIOR FILING DATE: 1998-01,02
NUMBER OF SEQ ID NOS: 122194;
COUNTAINS DATE: 1998-01,02
                                                       Query Match
Best Local Similarity
""" hes 7; Conserve
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; ORGANISM: Saccharomyces cerevisiae
US-09-335-032-2036
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LENGTH: 10
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GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Velculescu, Victor
APPLICANT: Vogelstein, Bert.
APPLICANT: Kinzler, Kenneth
TITLE OF INVENTION: Characterization of the Yeast
TITLE OF INVENTION: Transcriptome
FILE REFERENCE: 01107.78572
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1 tgttgga 7
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                                                                          Conservative
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100.0%; Pred. No.
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Query Match
Best Local Similarity
Thes 7; Conserve
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US-09-336-376-1209
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US-09-335-032-11545
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GENERAL INFORMATION:
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LENGTH: 10
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PRIOR APPLICATION NUMBER: US 60/035,917
PRIOR PILING DATE: 1997-01-23
PRIOR APPLICATION NUMBER: US 09/012,031
PRIOR PILING DATE: 1998-01-22
PRIOR FILING DATE: 1998-01-22
NUMBER OF SEQ ID NOS: 12219
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APPLICANT: Vogelstein, Bert
APPLICANT: Kinzler, Kenneth
APPLICANT: Kinzler, Kenneth
TITLE OF INVENTION: Characterization of the Yeast
TITLE OF INVENTION: Transcriptome
FILE REFERENCE: 01107.78572
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EARLIER APPLICATION NUMBER: 60/090,040
EARLIER FILING DATE: 1998-06-19
EARLIER APPLICATION NUMBER: 60/090,041
EARLIER FILING DATE: 1998-06-19
EARLIER APPLICATION NUMBER: 60/089,853
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT FILING DATE: 1999-01 EARLIER APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: POLYNUCLEOTIDE POPULATION ISOLATED TITLE OF INVENTION; HUMAN MONOCYTES
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                                                                                                           EARLIER
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                                                                    FILING DATE: 1998-06-19
APPLICATION NUMBER: 60/
FILING DATE: 1998-06-19
                                  APPLICATION NUMBER: 60/089,991 FILING DATE: 1998-06-19
                                                                                                                             APPLICATION NUMBER: 60/090,072
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                                                                                                                                                                                                                                            APPLICATION NUMBER: 60/090,035
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FILING DATE: 1998-06-19
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                APPLICATION NUMBER: 60/090,000
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1998-06-19
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Pred. No.
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EARLIER FILING DATE: 1998-12-08

NUMBER OF SEQ ID NOS: 5980

SOFTWARE: FASTSEQ for Windows Ver:
SEQ ID NO 1209

LENGTH: 10

TYPE: DNA

ORGANISM: Homo sapiens
US-09-336-376-1209
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Best Local Similarity
Thehes 7; Conserv
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                                CURRENT FILING DATE: 1999-06-17
EARLIER APPLICATION NUMBER: 60/090,039
EARLIER APPLICATION NUMBER: 60/090,039
EARLIER APPLICATION NUMBER: 60/090,040
EARLIER FILING DATE: 1998-06-19
EARLIER APPLICATION NUMBER: 60/090,041
EARLIER FILING DATE: 1998-06-19
EARLIER FILING DATE: 1998-06-19
EARLIER FILING DATE: 1998-06-19
EARLIER APPLICATION NUMBER: 60/089,853
EARLIER APPLICATION NUMBER: 60/089,997
EARLIER APPLICATION NUMBER: 60/090,079
EARLIER APPLICATION NUMBER: 60/090,079
EARLIER APPLICATION NUMBER: 60/090,079
EARLIER FILING DATE: 1998-06-19
EARLIER APPLICATION NUMBER: 60/090,079
EARLIER FILING DATE: 1998-06-19
EARLIER FILING DATE: 1998-06-19
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                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 4002,
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FILING DATE: 1998-06-19
APPLICATION NUMBER: 60/090,077
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FILING DATE: 1998-06-19
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EARLIER APPLICATION NUMBER: 60/090,048
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ER FILLING DATE: 1998-06-19
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ER FILLING DATE: 1998-06-19
ER APPLICATION NUMBER: 60/090,036
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ER APPLICATION NUMBER: 60/090,080
ER FILING DATE: 1998-06-19
ER APPLICATION NUMBER: 60/089,833
ER FILING DATE: 1998-06-19
ER APPLICATION NUMBER: 60/089,994
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FILING DATE: 1998-06-19
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DB seq length: 2000000000
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1: /cgn2_6/ptodata/1/pna/PCT_NEW_COMB.seq:*

2: /cgn2_6/ptodata/1/pna/US06_NEW_COMB.seq:*

3: /cgn2_6/ptodata/1/pna/US07_NEW_COMB.seq:*

4: /cgn2_6/ptodata/1/pna/US09_NEW_COMB.seq:*

5: /cgn2_6/ptodata/1/pna/US09_NEW_COMB.seq:*

6: /cgn2_6/ptodata/1/pna/US10_NEW_COMB.seq:*

7: /cgn2_6/ptodata/1/pna/US10_NEW_COMB.seq:*

8: /cgn2_6/ptodata/1/pna/US10_NEW_COMB.seq:*
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Match
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Copyright (c) 1993 - 2000 Comp
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US-09-825-805-181
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US-09-685-6648-1637
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US-09-740-332-3454
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1816, Appl
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   CURRENT APPLICATION NUMBER: US/09/825,805
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Sequence 624 Sequence 855 Sequence 116 Sequence 117 Sequence 108 Sequence 108 Sequence 70, Sequence 70, Sequence 72, Sequence 77, Sequence 709 Sequence 931 Sequence 539 Sequence 108
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PRIOR APPLICATION NUMBER: 09/578,223

2001-09-27

CURRENT FILING DATE:

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APPLICANT: Choi, Julie Y
APPLICANT: Choi, Julie Y
APPLICANT: Kliem, Stefanie E
APPLICANT: Kliem, Stefanie E
APPLICANT: Kliem, Stefanie E
APPLICANT: Ksehv, Beena
APPLICANT: Parks, Katie E
APPLICANT: Stephens, J. Claiborne
TITLE OF INVENTION: HAPLOTYDES OF THE TCF2 GENE
FILE REFERENCE: TCF2_MWH-1439PCT
CURRENT APPLICATION NUMBER: PCT/US01/30411
CURRENT FILING DATE: 2001-09-27
PRIOR APPLICATION NUMBER: 60/235,710
PRIOR APPLICATION NUMBER: 60/235,710
PRIOR APPLICATION NUMBER: 60/235,710
PRIOR FILING DATE: 2000-09-27
NUMBER OF SEQ ID NOS: 36
SOFTWARE: PatentIn version 3.1
SEQ ID NO 7
LENGTH: 15
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PRIOR APPLICATION NUMBER: 09/476,387
PRIOR APPLICATION NUMBER: 09/474,432
PRIOR APPLICATION NUMBER: 09/474,432
PRIOR APPLICATION NUMBER: 09/301.511
PRIOR FILING DATE: 1999-12-29
PRIOR APPLICATION NUMBER: 09/301.511
PRIOR FILING DATE: 1999-04-28
PRIOR APPLICATION NUMBER: 09/186,675
PRIOR APPLICATION NUMBER: 09/186,675
PRIOR FILING DATE: 1998-11-04
PRIOR FILING DATE: 1998-04-29
PRIOR APPLICATION NUMBER: 60/083,727
PRIOR APPLICATION NUMBER: 60/064,866
PRIOR FILING DATE: 1997-11-05
NUMBER OF SEQ ID NOS: 1558
SOFTWARE: PATENTIN VERSION 3.0
SEQ ID NO 181
LENGTH: 13
TYPE: RNA
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PCT-US01-30411-17/c
Sequence 17, Application PC/TUS0130411
GENERAL INFORMATION:
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Matches 4; Conservative
APPLICANT: Genaissance Pharmaceuticals, APPLICANT: Choi, Julie Y
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Вb

7 uguugga 13

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; SOFTWARE: PatentIn version 3
; SEQ ID NO 182
; LENGTH: 15
; TYPE: RNA
; ORGANISM: Homo sapiens
US-09-825-805-182
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APPLICANT: Stephens, J. Claiborne
TITLE OF INVENTION: HAPLOTYPES OF THE TCF2 GENE
FILE REBERENCE: TCF2_MWH-1439PCT
CURRENT APPLICATION NUMBER: PCT/US01/30411
CURRENT FILING DATE: 2001-09-27
PRIOR APPLICATION NUMBER: 60/235,710
PRIOR PILING DATE: 2000-09-27
NUMBER OF SEQ ID NOS: 36
SOFTWARE: Patentin version 3.1
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PCT-US01-30411-17
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Best Local Similarity
Matches 7; Conserv
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LENGTH: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Zinnen, Shawn
TITLE OF INVENTION: Nucleotide Triphosphate and their Incorporation into Oligonucl
FILLE REFERENCE: MBHB00.831-F (400/009)
CURRENT APPLICATION NUMBER: US/09/825,805
CURRENT APPLICATION NUMBER: US/09/825,805
PRIOR APPLICATION NUMBER: 09/476,387
PRIOR FILING DATE: 2000-05-23
PRIOR FILING DATE: 2000-12-30
PRIOR FILING DATE: 1999-12-30
PRIOR APPLICATION NUMBER: 09/474,432
PRIOR APPLICATION NUMBER: 09/474,432
PRIOR APPLICATION NUMBER: 09/474,432
PRIOR APPLICATION NUMBER: 09/301,511
PRIOR APPLICATION NUMBER: 09/31,511
PRIOR APPLICATION NUMBER: 09/31,511
PRIOR APPLICATION NUMBER: 09/31,511
PRIOR FILING DATE: 1999-10-28
PRIOR FILING DATE: 1999-10-28
                                             Matches
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Best Local Similarity
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PRIOR FILING DATE: 1998-11-04
PRIOR APPLICATION NUMBER: 60/083,727
PRIOR FILING DATE: 1998-04-29
PRIOR APPLICATION NUMBER: 60/064,866
PRIOR APPLICATION NUMBER: 60/064,866
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Beaudry, Amber
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Adamic, Jasenka Matulic
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Pred. No.
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Pred. No. 6.1e+04;
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US-09-527-030C-181

Sequence 181, Application US/09527030C

SEENERAL INFORMATION:

APPLICANT: VAN DOORN, Leen-Jan et al.

TITLE OF INVENTION: Detection and identification of Human Papillomavirus by PCR

TITLE OF INVENTION: Specific reverse hybridization.

FILE REFERENCE: 3501-0101P

CURRENT APPLICATION UNMBER: US/09/527,030C

CURRENT APPLICATION UNMBER: US/09/527,030C

CURRENT FILING DATE: 2000-03-16

NUMBER OF SEQ ID NOS: 497

SOFTWARE: PatentIn version 3.0
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Best Local Similarity
Thes 7; Conserv
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US-09-685-664B-1636
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Best Local Similarity
Thes 7; Conserv
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GENERAL INFORMATION:
APPLICANT: VAN DOORN, Leen-Jan et al.
TITLE OF INVENTION: Detection and identification of Human Papillomavirus by PCR and
TITLE OF INVENTION: Specific reverse hybridization.
FILE REFERENCE: 3501-0101P
CURRENT APPLICATION NUMBER: US/09/527,030C
CURRENT FILING DATE: 2000-03-16
NUMBER OF SEO ID NOS: 497
SOFTWARE: Patentin version 3.0
SEO ID NO 105
                                                                                                                                          Sequence 1636, Application US/09685664B GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQ ID NO 181
LENGTH: 17
                                                                             APPLICANT: Ribozyme Pharmaceuticals, APPLICANT: Pavco, Pam APPLICANT: McSwiggen, Jim
APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method and
TITLE OF INVENTION: Levels of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 16
TYPE: DNA
ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: Type specific probe derived from the Human Papillomavirus (HPV)
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100.0%; Pred. No. .6.1e+04;
Live 0; Mismatches 0;
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  Reagent for Treatment of Diseases or Conditions Relate Vascular Endothelial Growth Factor Recptor
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Pred. No.
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; TYPE: RNA; Homo sapiens; ORGANISM: Homo sapiens US-09-685-664B-1637
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                                                                                                                               US-09-685-664B-3484
               Sequence 3484, Application US/09685664B GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc. APPLICANT: Pavco, Pam APPLICANT: MCSWiggen, Jim
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Matches 4; Conserv
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Escobedo, Jaime TITLE OF INVENTION: Method and Reagent for Treatment of Diseases or Conditions Rel TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Recptor FILE REFERENCE: MBHB00-876-K (400/021) CURRENT APPLICATION NUMBER: US/09/685,664B CURRENT FILING DATE: 2000-10-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-10-26
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR FILING DATE: 1996-01-08
PRIOR APPLICATION NUMBER: US 09/371,772
PRIOR APPLICATION NUMBER: US 09/371,772
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APPLICANT: Pavco, Pam
APPLICANT: McSwidgen, Jim
APPLICANT: Stinchcomb, Dan
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APPLICANT:
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ORGANISM: Homo sapiens
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Stinchcomb,
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Pred. No. 6.1e+04;
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Pred. No. 6.1e+04;
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APPLICANT: McSwiggen, Jim Applicant: Escobedo, Jaine Escobedo, Jaine Filte OF INVENTION: Method and Reagent for Treatment of Diseases or Conditions Relate TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor FILE REFERENCE: Mathemol-876-K (400/021)

FILE REFERENCE: Mathemol-876-K (400/021)

CURRENT APPLICATION NUMBER: US 090-10-10

PRIOR APPLICATION NUMBER: US 090-584,040

PRIOR FILING DATE: 1995-10-26

PRIOR APPLICATION NUMBER: US 08/584,040

PRIOR FILING DATE: 1996-01-08

PRIOR FILING DATE: 1996-01-08

PRIOR PPLICATION NUMBER: US 09/371,772

PRIOR PRIOR DATE: 1999-08-10

SEQ ID NO 3485

LENGTH: 17
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TITLE OF INVENTION: Method and Reagent for Treatment of Diseases or Conditions Relate
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHB00-876-K (400/021)
CURRENT APPLICATION NUMBER: US/09/685,664B
CURRENT FILING DATE: 1905-10-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR FILING DATE: 1996-01-08
PRIOR FILING DATE: 1996-01-08
PRIOR FILING DATE: 1999-08-10
NUMBER OF SEQ ID NOS: 8231
SOFTWARE: Patentin version 3.0
SEQ ID NO 3484
LENGTH: 17
В
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Best Local Similarity
Thes 4; Conserv
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US-09-685-664B-3485
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ORGANISM: Mus musculus US-09-685-664B-3486
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; Sequence 3493, Application US/09685664B
; GENERAL INFORMATION:
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TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Recptor
FILE REFERENCE: MBHB00-876-K (400/021)
FILE REFERENCE: MBHB00-876-K (400/021)
FILE REPLICATION NUMBER: US/09/685,664B
CURRENT FILING DATE: 2000-10-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-10-26
PRIOR FILING DATE: 1996-01-08
PRIOR FILING DATE: 1996-01-08
PRIOR FILING DATE: 1996-01-08
PRIOR FILING DATE: 1996-08-10
NUMBER OF SEQ ID NOS: 8231
SOFTMARE: Patentin version 3.0
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SOFTWARE: PatentIn version 3.0
SEQ ID NO 3493
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Ribozyme Pharmaceuticals, Inc.
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PRIOR APPLICATION NUMBER: US 09/371,772
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GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate
TITLE OF INVENTION: Hepatitis C Virus Infection
FILE REFERENCE: RPI 400/003
CURRENT FILING DATE: 2001-03-26
NUMBER OF SED ID NOS: 9704
SOFTWARE: Patentin version 3.0
SED ID NO 1100
LENGTH: 17
TYPE: RNA
ORGANISM: artificial sequence
FEATURE:
NAMEJKEY: misc_feature
COCATION:
OTHER INFORMATION: oligonucleotide substrate
US-09-740-332-1100
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NAME/KEY: misc_feature
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OTHER INFORMATION: oligonucleotide substrate
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Search completed: December 15, 2001, 05:58:20 Job time: 12263 sec
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Sequence 1101, Application US/09740332

GENERAL INFORMATION:

APPLICANT: Ribozyme Pharmaceuticals Inc.

TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate
TITLE OF INVENTION: Hepatitis C Virus Infection

FILE REFERENCE: RPI 400/003

CURRENT APPLICATION NUMBER: US/09/740,332

CURRENT FILING DATE: 2001-03-26

NUMBER OF SEQ ID NOS: 9704

SOPTWARE: PATENTIN VERSION 3.0

SEQ ID NO 1101

LENGTH: 17

TYPE: RNA

ORGANISM: artificial sequence
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Matches 7; Conservative
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SUMMARIE

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ALIGNMENTS

RESULT 1
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LEPtospira fainei 16S ribosomal RNA gene, partial sequence.
ACCESSION
U60594.1 GI:1408219
KEYWORDS
SOURCE
ORGANISM
REFERENCE
AUTHORS
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TITLE
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AUTHORS
Billinghurst,M.L., Letocart,M., Merien,F. and Serrano,M.S.
Billinghurst,M.D., Baranton,G., Bulach,D.M.,
Billinghurst,M.L., Letocart,M., Merien,F., Serrano,M.S.
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Leptospira fainei par
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Y19243.1 GI:6318184
16S ribosomal RNA; 16
                                              2 (bases 1 to 1450)
Krogfelt,K.A.
Direct Submission
Submitted (05-NOV-1999) K.A. Krogfelt,
Dept. of Gastrointestinal Infections, A
Copenhagen, DENMARK
                                                                                                Unpublished
2 (bases 1
                                                                                                                 Petersen, A.M., Krogfelt, K.A., Schlichting, P.
Leptospira fainel serovar Hurr
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/organism="Leptospira
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/db_xref="taxon:48782"
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Nucleic P
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Submitted (22-NOV-1989) Fukunaga M., Faculty
Pharmaceutical Science, Fukuyama University,
Fukuyamashi, Hiroshimaken 729-0, Japan
2 (bases 1 to 1508)
Fukunasa W 1971
                                                                                                                        Fukunaga, M., Horite, I., Okuzako, N.
Nucleotide sequence of a 16S rRNA
serovar canicola strain Moulton
Nucleic Acids Res. 18 (2), 366 (19
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Best Local Sim
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                                   L.inadai g
Z21634
Z21634.1
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Direct Submission

Submitted (09-FEB-1993) Hookey J.V., Public Health Laboratory

Service, Leptospira Reference Laboratory, Stonebow Road, Hereford,

Herefordshire, United Kingdom, HR1 2ER
                                                                                                                                                                                                                                                                       Bacteria; Spirochaetales;
1 (bases 1 to 1374)
Hookey,J.V.
                                                                                                                                                                                                                                                       Phylogeny of Leptospiraceae Unpublished
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Leptospira inadai
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                                                                        Conservative
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/strain="LYME"
/db_xref="taxon:29506"
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/product="16S ribosomal RNA"
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                                                                     aagatttattgcttggagatgagcccgcggccgattagctagttggtgaggtaatggctc
cggtccatactcctacgggaggcagcagttaagaatcttgctcaatgggggaaaccctga
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                                                             AAGATTTATTGCTCGGAGCTGAGCCCGCGCCCGATTAGCTAGTTGGTGAGGTAATGGCTC
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U12676.1
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Direct Submission
Submitted (25-JUL-1994) Michael
Biological Research, 11099 North
92037, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1483 bp DNA Leptospira weilii Cellidoni seguence.
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Phylogenetic evidence for horizontal transfer sequence between species in a spirochete genus J. Bacteriol. 176 (19), 5982-5987 (1994)
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Leptospira weilii
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/strain="serogroup Celledoni,
Celledoni (ATCC 43285)"
/isolate="WA45"
                                                                                                                                                                                                                                                                                      /product="16S rRNA"
331 c 452 g
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94.1%;
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North Torrey Pines Road, La
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1 Leptospira weilii W

U12677

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ralph,D. and McClelland,M.
Phylogenetic evidence for horizontal transfer of sequence between species in a spirochete genus
J. Bacteriol. (1994) In press
2. (bases 1 to 1489)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Direct Submission
Submitted (25-UUL-1994) Michael McClelland, California
Biological Research, 11099 North Torrey Pines Road, La
92037, USA
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Bacteria; Spirochaetales; Leptospiraceae;
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/strain="serogroup Hebdomadis, subserogroup
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Leptospira borgpetersenii.

M Leptospira borgpetersenii

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Leptospira borgpetersenii

1 (bases 1 to 1513)

M Leptospira borgpetersenii

Ralph, D. and McClelland, M.

Phylogenetic evidence for horizontal tr

sequence between species in a spirochet

J. Bacteriol. (1994) In press

2 (bases 1 to 1513)
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Direct Submission
Submitted (25-JUL-1994) Michael
Biological Research, 11099 North
92037, USA
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/strain="serovar hardjo, strain Hardjo
/isolate="1950"
/db_xref="taxon:174"
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Ralph,D. and McClelland,M.
Phylogenetic evidence for horizontal transfer c
sequence between species in a spirochete genus
J. Bacteriol. 176 (19), 5982-5987 (1994)
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Leptospira borgpetersenii
Bacteria: Spirochaetales; Leptospiraceae; Leptospira
1 (bases 1 to 1489)
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Submitted (25-JUL-1994) Michael N
Biological Research, 11099 North
92037, USA
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/strain="serovar balcanica, strain 16;
/isolate="BD30"
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Leptospira noguchi i
U12671
U12671.1 GI:558930
Bacteria; Spirochaetales;
1 (bases 1 to 1516)
Ralph,D. and McClelland,M
Phylogenetic evidence for
                      Leptospira noguchii.
Leptospira noguchii
                                       GI:558930
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Submitted (25-JUL-1994) Michael M
Biological Research, 11099 North
92037, USA
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J. Bacteriol. 176
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/organism="Leptospira noguchii"
/strain="serogroup Autumnalis,
serover fortbragg, strain Fort
/isolate="NB36"
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Hookey, J.V.
Direct Submission
Submitted (09-FEB-1993) Hookey
Service, Leptospira Reference I
                                                                 Leptospira borgpetersenii.
Leptospira borgpetersenii
Bacteria; Spirochaetales;
1 (bases 1 to 1425)
                                                                                                   Z21630.1 GI:433579
16S ribosomal RNA.
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/strain="JAVANICA VELDRAT BATAVIA 46"
/db_xref="taxon:174"
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                                                                            McClelland,M.
Direct Submission
Submitted (25-JUL-199)
Biological Research,
92037, USA
                                                                                                                             Ralph, D. and McCielland, M. Phylogenetic evidence for horizontal transfer sequence between species in a spirochete genus J. Bacteriol. 176 (19), 5982-5987 (1994)
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Leptospira weilii
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Sarmin, serovar sarmin, strain Sarmin"
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/db_xref="taxon:28184"
     /product="16S rRNA"
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l Research, 11099 North Torrey Pines Road, La
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Leptospira kirschneri.
Leptospira kirschneri
Bacteria; Spirochaetales; L
1 (bases 1 to 1419)
Hookey,J.V.
                                                                                                                                     Direct Submission
Submitted (09-FEB-1993) Hookey J.V., Public Health
Service, Leptospira Reference Laboratory, Stonebow
Herefordshire, United Kingdom, HRI 2ER
                                                                                                                                                                                                                                              L.kirschneri gene fo
Z21628
Z21628.1 GI:433582
                                                                                                                                                                     Hookey, J.V.
                                                                                                                                                                                  Unpublished
                                                                                                                                                                                           Phylogeny of Leptospiraceae
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                                     Similarity
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nilarity 94.2%;
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                                                                         /organism="Leptospira kirschneri"
/strain="CYNOPTERI 3522 C"
/db_xrei="taxon:29507"
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/product="16S ribosomal RNA"
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Submitted (25-JUL-1994) Michael McClelland, California
Biological Research, 11099 North Torrey Pines Road, La
92037, USA
                                                                                                                                                                                                                                                                                                                                                                                 LSU12672 1494 bp
Leptospira santarosai
U12672
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Phylogenetic evidence for horizontal transfer of sequence between species in a spirochete genus J. Bacteriol. 176 (19), 5982-5987 (1994)
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Leptospira santarosai
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                                                                                                                   Conservative
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a 332 c 453 g 311 t 15 others
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93.2%;
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agctaataccggatagtcctgttggatcacaagattttgataggtaaagatttattgcttg 194
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L.meyeri gene for 16S ribosomal RNA (partial).
Z21648
Z21648.1 GI:433585
                                                                                                                                                                                                                                                                                                                                                                                                                    2 (bases 1 to 1418)
Hookey, J.V.
Direct Submission
Submitted (09-FEB-1993) Hookey J.V., Public Health Laboratory
Service, Leptospira Reference Laboratory, Stonebow Road, Hereford,
Service, Inted Kingdom, HRI 2ER
Location/Qualifiers
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Leptospira meyeri
Bacteria; Spirochaetales; Leptospiraceae; Leptospira.
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324 c 421 g 299 t
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cc 1332 	273 accccatgaagtcggaatcgctagtaatcgcggatcagcatgccgcggttgaatacgttc 	Qy 12 Db 12	
cg 1272 CG 1259	213 togoaagagggagctaatototaaaagtoggtoocagttoggattggggtotgcaacto 	Qy 12 Db 12	
ac 1212 	53 atggcctttatgtccagggccacacacgtgctacaatggccgatacagagggtcgccaa 	Qy 11 Db 11	
tc 1152	93 gcactggtacgaaactgccggtgacaaaccggaggaaggcggggatgacgtcaaatcc 	Qy 10 Db 10	
39 1092 11 36 1079)34 tgttgggttaagtcccgcaacgagcgcaacccctatcgtatgttgctacc-ttaagttg 	Qy 10	
ja 1033 SA 1019	774 tatgageettegggeagatteaeaggtgetgeatggttgtegteagetegtgtegtgag 	Qy 9	
ia 973 A 959	114 ttcgatgataccccaaaaacctcacctgggcttgacatggatctgaatcatgtagagat 	ОУ 9 Db 9	
la 913 A 899	54 aagagtgaaactcaaaggaattgacgggggtccgcacaagcggtggagcatgtggttta 	Qy 8 Db 8	
C 853	4 gttttaaccctcagtaacgaacctaacggattaagtagaccgcctggggactatgctcg 	Qy 7 Db 7	
9 793 G 779	34 taaacgggattagataccccggtaatccacgccctaaacgttgtctaccagttgttggg 	Qy 7 Db 7	
9 733 G 719	74 ccagtggcgaaggcgacttgctggctcaaaactgacgctgaggcacgaaagcgtgggta 	Qy 6	
a 673 A 659	14 tgggagaggcaagtggaattccaggtgtagcggtgaaatgcgtagatatctggaggaac 	Qу 6 Db 6	
t 613 T 599	54 caggigigaaaacigcgggcicaacccgiggccigcactigaaaciacaagiciggagi 	Qy 5	
t 553 539	94 atggtgcaagcgttgttcggaatcattgggcgtaaagggtgcgtaaggcggatttgtaag 		
t 493 479	4 tgatgatggtacctgcctaaagcaccggctaactacgtgccagcagccgcggtaatacg	0y 4	
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Direct Submission

Submitted (19-JUN-1992) Hookey J. V., Public Health Laboratory Service, Leptospira Reference Laboratory, Stonebow Road, Herefordshire, United Kingdom, HRI 2ER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bacteria; Spirochaetales; Leptospiraceae; Leptospira.

1 (bases 1 to 1396)
Bryden, J., Gatehouse, L., Gatehouse, J.A. and Hookey, J.V.
Automated PCR cycle sequencing of 16S ribosomal RNA genes
Unpublished
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/strain="icterchaemorrhagiae RGA"
/db_xref="taxon:173"
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                               New isolated pathogenic Leptospira bacterium - useful for, developing products for conferring protective immunity, and prophylactic or therapeutic treatment
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     an organism in various of environments, especially polluted environments (particularly those contaminated with oil or harmful chemicals) using migratory analysis. The methods enable the detection and quantitation of a microorganism obtained from a polluted site in order to identify strains capable of degrading petroleum and harmful chemicals. Such organisms are particularly used for environmental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Detection and quantitation of microorganism having specific function and its gene from natural environment, with identification of petroleum-digesting bacterium cycloclasticus for treating oil spillage
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                                                                                                                                                                                                                                                                                                                                                                           decontamination. The invention is particularly concerned with the detection and quantitation of petroleum-degrading bacteria of the genus Cycloclasticus. Cycloclasticus pugetii 16S rRNA gene sequences (AAF76234-AAF76237) are specifically claimed, as are probes (AAF76238-AAF76240) which hybridise to these sequences. The present sequence is one of the claimed Cycloclasticus pugetii 16S rRNA (ribosomal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (AGEN ) AGENCY OF IND SCI & TECHNOLOGY.
(NISH-) NISHIMATSU CONSTRUCTION CO LTD.
(NYKL-) NYK LOGISTICS TECHNOLOGY INST CO LTD.
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hydrocarbon; detection; quantitation; identification; oil spill;
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                                                                                                                                                                                                                                                                                                                    Sequence 1528 BP; 394 A; 343 C; 467 G; 323 T; 1 other;
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                                     68 atgctagcttgctagcaggcgtcgagtggcggacgggtgagtaatgcataggaatctacc 127
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509	agcaccggctaactacgtgccagcagccgcggtaatacgtatggtgcaagc	Qy
485	ttcagtagggaggaaagtttaaggttaataaccttaggccctgacgttacctaca	Db
452	aggcaggaaaaataagcagcaatgtgatgatggt	Qy
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                           The invention relates to a method for detecting, identifying and quantitating a microorganism which has a specific function, or DNA therefrom, from the environment. The method comprises broadly classifying the microorganism after serial dilution; amplifying and cloning specific gene domains using extracted DNA as a template; examining the differences between the gene domains cloned and determining their base sequences; and positively identifying the microorganism from the base sequence data. The invention also relates to a method to evaluate the function of such an organism in various of environments, especially polluted environments (particularly those contaminated with oil or harmful enventions of a microorganism obtained from a polluted site in order to identify strains capable of degrading petroleum and harmful chemicals. Such organisms are particularly used for environmental deportance of the particular of the p
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Best Local Similarity
Matches 1137; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Monitoring of oil contamination of sea water - where oil contamination is evaluated by detection of a microbe having properties from e.g. having no flagellum, being a Gram-negative bacterium, belonging to Proteobacteria, gamma subdivision, etc.z
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The sequences given in AAC86021-30 represent 16s rDNA sequences from CC polyphosphate accumulating organisms (PAOs). Sequences which are unique CC to these 16s rDNA sequences are used to create a probe or primer for CC detecting the relevant organisms. The primer/probe sequences are CC useful for detecting pAO cells in a sample, by treating cells in the CC sample to fix cellular contents, contacting fixed cells with the CC sample, under conditions which allow the probe to hybridize with 16s CC hapten, under conditions which allow the probe to hybridize with 16s CC rRNA within the fixed cell, removing unhybridized probe from the fixed CC cells, and detecting the labeled probe-NNA hybrid by fluorescence in CC situ hybridization. The primer/probe sequences are useful for CC identifying pAOs that are capable of biologically removing compositions from waste water. Rapid assessment of the presence of a phosphorus from waste water sample, can be done using the primer/CC probe sequences. They allow quick and convenient assessment of whether CC a sludge or waste water sample includes PAOs and allows quantitation of PAOs cells in samples.
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RESULT 8
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Best Local Similarity
Matches 1133; Conser The invention relates to a new method for the production of taxane and pacifitaxel compounds that comprises culturing bacteria isolated from a plant species of Taxus or bacteria produced by mutating the isolated bacteria. The methods can be used for the production of taxanes such as paclitaxel, 10-deacetylcephalomannine, 7-epitaxol, 10-deacetyl-7-epitaxol, 7-epicephalomannine, baccatin III, 710-deacetyl-baccatin III, cephalomannine, 7-epibaccatin III, 7-xylosyltaxol, 7-xylosyl-cephalomannine, taxagifine, delta-benzoyloxy taxagifine, 9-acetyloxy taxusin, 9-hydroxy taxusin, taxane ID, taxane ID, taxane IC, or taxane ID (claimed). The taxane and paclitaxel compounds can be used for the treatment of cancers. The methods can provide for the production of taxanes and paclitaxel at a concentration of 1-25 mu g/l. Sequence 1512 Disclosure; Fig 8F; 93pp; WPI; 1999-418940/35 Page Boissinot M, WO9932651-A1 Bacillus subtilis 10-SEP-1999 22-DEC-1997; 18-DEC-1998; (BCMB-) BCM DEV 308 248 219 188 166 128 106 83 56 subtilis partial 16S rRNA 8 ب agatttgataggtaaag-----atttattgcttggagatgagcccgcgggccgattagc gatcatggctcagaactaacgctggcggcgcgtcttaaacatgcaagtcgagcgg----ggccacaatggaactgagacacggtccatactcctacgggaggcagcagttaagaatctt | tagttggtgaggtaatggctcaccaaggcgaccgatcggtagccggcctgagagggtgtcc gtaagactgggataactccgggaaatcggggctaataccggatggttgtttgaaccgcat ccgagtctgggataactttccgaaaggaaagctaataccggatagtcctgttggatcaca tagttggtgaggtaacggctcaccaaggcgacgatgcgtagccgacctgagagggtgatc ggttcagacataaaaggtggcttcggctaccacttacagatggacccgcggcgcattagc tgggagcttgctccctgatgttagcggcggacgggtgagtaacacgtgggtaacctgcct 127 gatcctggctcaggacgaacgctggcggcgtgcctaatacatgcaagtcgagcgaacaga paclitaxel; plant; bacteria; of taxane and Conservative (first ВP; Gagne M, 97US-0995960 98WO-CA01150 377 A; 353 C; 480 entry) 55.18; 75.48; Harvey paclitaxel English. 0; Score 813.4; DB 20; Pred. No. 8.6e-243; 0; Mismatches 326; gene Ķ sequence Helie G; compounds mutation; 302 ζ. T; 0 other; Landry N; cancer; 16S rRNA; Indels Length 1512; 44; Gaps 187 165 105 67 367 307 278 247 218 55

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Query Match
Best Local Similarity
Matches 921; Conserv

Conservative

220;

Mismatches

54.98;

Score 810.2; Pred. No. 8.

.5e-DB

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Gaps

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                                                                                                This sequence encodes a Bacillus species pectate lyase of the invention. The pectate lyase enzymes are obtained from Bacillus halodurans (BH) agaradhaerens (BH), Bacillus licheniformis (BL), Bacillus halodurans (BH and other Bacillus species. The pectate lyase enzymes can be used for degrading pectin, pectate and polygalacturonides. They can be used in detergent compositions, for cleaning hard surfaces, for machine treatmen of fabrics, for improving the properties of cellulosic fibres, yarn, woven or non-woven fabric, for the degradation of plant material (e.g. recycled waste paper, mechanical paper-making pulps or fibres subjected to a retting process, for preparing animal feed or for processing wine or juice. DNA encoding the enzymes can also be used for the production or transcent.
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24-NOV-1997;
06-MAY-1998;
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                                                                 10-JUN-1998;
20-OCT-1998;
20-OCT-1998;
28-OCT-1998;
28-OCT-1998;
23-DEC-1998;
05-MAR-1999;
05-MAR-1999;
05-MAR-1999;
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11-MAR-1999;
11-MAR-1999;
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synthetic fibre; yarn; fabric; printing paste; thickener; viscosity;
plant material degradation; recycled waste paper; paper making pulp;
guar; locust bean gum; mannan-containing food; coffee extract;
cleaning composition; machine washing; hard-surface cleaner; 16S RNA
dishwashing; oral; dental; contact lens; body-care composition;
fabric softener; oil well drilling; subterranean formation fracture;
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endo-1,4-mannanase; galactomannan; 1,4-beta-D-mannosidic
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The mannanase hydrolyses galactomannans. Specifically, mannanases CC hydrolyse 1,4-beta-D-mannosidic linkages in mannans, galactomannans, and galactomannans. The mannanase protein, or CC glucomannans, and galactoglucomannans. The mannanase protein, or CC preparations containing it, are used to improve properties of cellulosic CC or synthetic fibres, yarn or (non)woven fabrics (removal of mannan-based CC sizes or printing pastes). They are also used to degrade or modify CC plant materials (particularly recycled waste paper, paper making pulps, CC or material containing guar or locust bean gums (thickeners), or to CC reduce viscosity of mannan-containing foods or feeds). The mannanases CC are also used to process coffee extracts (to inhibit gel formation); in CC cleaning compositions (for machine washing of fabrics, as hard-surface CC cleaners, for hand or machine dishwashing, also in oral, dental, contact CC cleaners, for by care compositions) where they remove mannan-containing CC solls and prevent binding of some solls to cellulosics; and in fabric CC subterarean formations.
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Gram-negative bacterium;
alkane; 16S rDNA gene; ds
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cgggaggacgttcaccacggtgtggttcatgactggggtgaagtcgtaacaaggtagc
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RESULT 13
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                                                                                                                                                                                                                                                                                   Sequence
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Best Local Similarity 76.1%;
Matches 1140; Conservative
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US-08-95-960-2
US-08-95-953-160
US-09-198-955A-13
US-09-549-1108-2
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US-09-198-955A-14
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ALIGNMENTS

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REFERENCE/DOCKET NUMBER: UOFW
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 682 8100
TELEFAX: (206) 224 0779
TELEX: 4938023
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENCTH: 1536 base pairs
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Patent No. 5874291
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: Christensen O'Connor JG
STREET: 1420 Fifth Avenue, Suite 1
CITY: Seattle
STATE: Washington
COUNTRY: USA
ZIP: WA 98101-2347
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
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FILING DATE:
FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/
FILING DATE: 20-MAX-1994
ATTORNEY/AGENT INFORMATION:
NAME: Sheiness, Diana K.
REGISTRATION NUMBER: 35,35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOCTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
                   HYPOTHETICAL:
ANTI-SENSE: NO
                                                     MOLECULE TYPE: nucleic acid
DESCRIPTION: "16S ribosoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: Degradation of Environmental TITLE OF INVENTION: Filamentous Bacterium
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
                                                                                                      TYPE: nucleic acid
STRANDEDNESS: sing
TOPOLOGY: linear
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1420 Fifth Avenue, Suite 2800
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Bielefeldt, Angela
Gransel, H. David
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Strand, Stuart
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Matches 1113; Conservative
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                         GACGGGGACCCGCACAAGCGGTGGATGATGTGGTTTAATTCGATGCAACGCGAAAAAACCT
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cacctgggcttgacatggatctgaatcatgtagagatatatgagccttcgggcaga----
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Mismatches 343;
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US-09-198-955A-14
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           APPLICANT:
APPLICANT:
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                                                                                                       INFORMATION:
Glad, Sanne O. S.
Kauppinen, Markus
Schnorr, Kirk
                                                     Bjornvad, Mads E.
                                                                Andersen, Lene N
Schulein, Martin
Lange, Niels E.
                                        Soren
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US-09-108-955A-14

; Sequence 14, Application US/09198955A
; Patent No. 6187580

GENERAL INFORMATION:
APPLICANT: Andersen, Lene N.
APPLICANT: Schulein, Martin
APPLICANT: Honge, Niels E.
APPLICANT: Honge, Niels E.
APPLICANT: Honge, Niels E.
APPLICANT: Moller, Soren
APPLICANT: Moller, Soren
APPLICANT: Kauppinen, Markus S.
APPLICANT: Kongsbak, Lars
TITLE OF INVENTION: No. 6187580el Pectate Lyases
FILE REFERENCE: 5378.200-US
CURRENT FILING DATE: 1998-11-24
PRIOR APPLICATION NUMBER: 1333/97
PRIOR FILING DATE: 1997-11-24
PRIOR APPLICATION NUMBER: 1344/97
PRIOR APPLICATION NUMBER: 60/67,249
PRIOR APPLICATION NUMBER: 60/67,249
PRIOR APPLICATION NUMBER: 09/073,684
PRIOR APPLICATION NUMBER: 09/073,684
PRIOR FILING DATE: 1998-11-02
PRIOR APPLICATION NUMBER: 09/073,684
PRIOR FILING DATE: 1998-05-06
PRIOR FILING DATE: 1998-11-02
INUMBER OF SEQ ID NOS: 32
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 14
LENGTH: 1508
TYPE: RNA
ORGANISM: Bacillus Sp.

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RESULT 3
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APPLICANT: Van Sol
APPLICANT: William
APPLICANT: IVETSON
APPLICANT: Farrell
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Patent No. 6140095
STREET: 2000 Pennsylvania Avenue, NW CITY: Washington STATE: DC COUNTRY: USA ZIP: 20006-1812 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk COMPUTER: 1BM PC COMPATIBLE US/08/501,126 FILING DATE: 29-DEC-1995 CLASSIFICATION NUMBER: US/08/501,126 FILING DATE: 29-DEC-1995 CLASSIFICATION NUMBER: 29.959 REFERENCE/DOCKET NUMBER: 4615-0057.00 TELECOMMUNICATION INFORMATION: TELEPHONE: (202) 887-1500
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Van Der Kleij, Wilhelmus
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Farrell, Roberta L.
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SEQUENCE CHARACTERISTICS:
LENGTH: 1521 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
HYPOTHETICAL: NO
ORIGINAL SOURCE:
ORGANISM: Bacillus Sp.
INDIVIDUAL ISOLATE: DSM 87
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Best Local Similarity 76.9%;
Matches 1097; Conservative
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RESULT 4 US-08-114-695A-7 Sequence 7, Application US/08114695A Patent No. 5508193 COUNTRY: USA
ZIP: 55402
ZUP: 55402
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, V.
CURRENT APPLICATION DATA: GENERAL INFORMATION: APPLICATION NUMBER: US/08, FILING DATE: 31-AUG-1993 CLASSIFICATION: 435 ATTORNEY/AGENT INFORMATION: NAME: MUETING, ANN M. REGISTRATION NUMBER: 33,97 APPLICANT: Mandelbaun
APPLICANT: Wackett, I
TITLE OF INVENTION: I
TITLE OF INVENTION: V
NUMBER OF SEQUENCES: CORRESPONDENCE ADDRESS: ADDRESSEE: SCHWEGMAN, LUNDBERG STREET: 3500 IDS CENTER CITY: MINNEAPOLIS STATE: MN REFERENCE/DOCKET NUMBER: Mandelbaum, Raphael WATER 8 DEGRADATION OF Lawrence US/08/114,695A 33,977 600.268USI Ŧ. & WOESSNER, S-TRIAZINES Version #, סי H TIOS

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TELEPHONE: 612-339-0331
TELEFAX: 612-339-3061
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 1536 base pairs
TYPE: nucleic acid
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Matches 897;
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ORIGINAL SOURCE:
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TOPOLOGY: linear
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                                 UGGGCCUGCACUGACGCUCAUGCACGAAAGCGUGGGGAGCAAACAGGAUUAGAUACCCUG
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Pred. No. 3e-280;
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CGCUUACCACGGGGGUUCGUGACUGGGGGUGAAGUCGUAACAAGGUAGC
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RESULT 5
US-08-995-960-2
"Coruence 2, Ar Sequence 2, Application Patent No. 6030818 GENERAL INFORMATION: CORRESPONDENCE ADDRESS:
ADDRESSEE: Dike, Bron.
STREET: 130 Water Str.
CITY: Boston
STATE: MA
COUNTRY: U.S.A.
ZIP: 02109 APPLICANT: PAGI, Michel APPLICANT: LANDRY, Natha. APPLICANT: BOISSINOT, Maried APPLICANT: HILLE, Maried APPLICANT: HARVEY, Mario APPLICANT: GAGNI, Martin APPLICANT: GAGNI, Martin APPLICANT: HARVEY, Mario
APPLICANT: GAGNI, Martin
TITLE OF INVENTION: BACTERIAL MASS
TITLE OF INVENTION: OF TAXANES AND
NUMBER OF SEQUENCES: 18 COMPUTER READABLE FORM: MEDIUM TYPE: Diskette COMPUTER: IBM Compatible OPERATING SYSTEM: Windows SOFTWARE: FastSEQ for Win LANDRY, Nathalie BOISSINOT, Maurice HILIE, Marie-Claude FastSEQ for Windows Bronstein, Street us/08995960 Roberts Version PRODUCTION PACLITAXEL δı Cushman, 2.0b

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US-08-995-960-2
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Best Local Similarity 75.3%;
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS
FENGTH: 1512 base pair
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APPLICATION NUMBER: US/08/995,96
FILING DATE: 22-DEC-1997
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: RESNICK, David S
REGISTRATION NUMBER: 34,235
REFERENCE/DOCKET NUMBER: 1826/47
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
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STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: rRNA
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Pred. No. 1.6e-279;
0; Mismatches 322;
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; Sequence 160, Applicate
; Patent No. 5843669
; GENERAL INFORMATION:
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APPLICANT: Kaiser, Michael W.
APPLICANT: Lyamichev, Victor I.
APPLICANT: Lyamichev, Natasha
TITLE OF INVENTION: Cleavage Of Nucleic i
TITLE OF INVENTION: Thermostable FEN-1 ENUMBER OF SEQUENCES: 190
CORRESPONDENCE ADDRESS:
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NAME: Ingolia, Diane E.
REGISTRATION NUMBER: 40,027
REFERENCE/DOCKET NUMBER: FORS-
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 705-8410
TELEPHONE: (415) 397-83381
INFORMATION FOR SEQ ID NO: 160:
SEQUENCE CHARACTERISTICS:
LENGTH: 1555 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
US-08-757-653-160
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 54.3%;
Best Local Similarity 75.3%;
Matches 1134; Conservative
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
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FILING DATE:
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: TOTALL NAME: F
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STREET: 220 Montgomery Street, St
CITY: San Francisco
STATE: California
COUNTRY: United States Of America
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ed. No. 1.9e-279;
Mismatches 324;
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RESULT 7
US-09-198-955A-13
; Sequence 13, Application
; Patent No. 6187580
; GENERAL INFORMATION:
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FILE CE ANNOUNCE: 5378.200-US
FILE REFERENCE: 5378.200-US
CURRENT FILING DATE: 1998-11-24
PRIOR APPLICATION NUMBER: 1134/97
PRIOR APPLICATION NUMBER: 1344/97
PRIOR FILING DATE: 1997-11-24
PRIOR APPLICATION NUMBER: 60/067,249
PRIOR APPLICATION NUMBER: 60/067,249
PRIOR FILING DATE: 1997-12-02
PRIOR FILING DATE: 1997-12-02
PRIOR FILING DATE: 1997-12-02
PRIOR FILING DATE: 1997-12-03
PRIOR FILING DATE: 1997-13-03
PRIOR FILING DATE: 1998-05-06
PRIOR APPLICATION NUMBER: 09/073,684
PRIOR APPLICATION NUMBER: 09/073,684
PRIOR FILING DATE: 1998-05-06
PRIOR APPLICATION NUMBER: 09/184,217
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SOFTWARE: FastSEQ for
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Best Local Similarity
Matches 916; Conserv
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ORGANISM: Bacillus
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Bjornvad, Mara-
Molle-
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RESULT 8
US-09-248-528-2
IS-09-248-528-2
; Sequence 2, Application US/09248528C
; Patent No. 6153415
; GENERAL INFORMATION:
; APPLICANT: Oriel, Patrick J
; APPLICANT: Padmakumar, Rugmini

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TITLE OF INVENTION: Hydratase from a Thermophilic Bacillus
TITLE OF INVENTION: Hydratase from a Thermophilic Bacillus
FILE REFERENCE: MSU 4.1-401
CURRENT FILING DATE: 1999-02-10
EARLIER APPLICATION NUMBER: US/09/248,528C
CURRENT FILING DATE: 1999-02-10
EARLIER APPLICATION NUMBER: 60/083,485
EARLIER FILING DATE: 1998-04-29
NUMBER OF SEQ ID NOS: 18
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
Matches 1128; Conserv
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NAME/KEY: YRNA
LOCATION: (1)..(1517)

PUBLICATION INFORMATION:

DATABASE ACCESSION NUMBER: Z26929

DATABASE ENTRY DATE: 1998-07-02
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Pred. No. 4e-276;
0; Mismatches 326;
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RESULT 9
US-09-549-108-2
US-09-549-108-2
; Sequence 2, Application US/09549108
; Patent No. 6214603
; Patent No. 6214603
; Patent INFORMATION:
APPLICANT: Padmakumar, Rugmini
APPLICANT: Rim, Sang H
TITLE OF INVENTION: Method for Producing Am
TITLE OF INVENTION: Hydratase from a Thermo
FILE REFERENCE: MSU 4.1-486
; CURRENT APPLICATION NUMBER: US/09/549,108
; CURRENT FILING DATE: 2000-04-13
; PRIOR APPLICATION NUMBER: 60/083,485
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PRIOR APPLICATION NUMBER: 09/24:
PRIOR FILING DATE: 1999-02-10
NUMBER OF SEQ ID NOS: 18
SOFTWARE: Patentin Ver. 2.0
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NAME/KEY: rRNA
LOCATION: (1)..(1517)
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DATABASE ENTRY DATE: 1998-07-02
US-09-549-108-2
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Best Local Similarity 75.0%;
Matches 1128; Conservative
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Pred. No. 4e-276;
0; Mismatches 326;
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GENERAL INFORMATION:

APPLICANT: Oriel, Patrick J

APPLICANT: Padmakumar, Rugmini

APPLICANT: Kim, Sang H

TITLE OF INVENTION: Method for Producing Amide Compounds Using

TITLE OF INVENTION: Hydratase from a Thermophilic Bacillus

FILE REFERENCE: MSU 4.1-489

CURRENT APPLICATION NUMBER: US/09/549,111

CURRENT FILING DATE: 1000-04-13

PRIOR APPLICATION NUMBER: 60/083,485

PRIOR APPLICATION NUMBER: 09/248,528

PRIOR APPLICATION NUMBER: 09/248,528

PRIOR APPLICATION NUMBER: 09/248,528

PRIOR FILING DATE: 1999-02-10

NUMBER OF SEQ ID NOS: 18

SOFTWARE: Patentin Ver. 2.0

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FEATURE:
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DATABASE ACCESSION NUMBER: Z26929
DATABASE ENTRY DATE: 1998-07-02
IS-09-549-111-2
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APPLICANT: Kim, Sang H

TITLE OF INVENTION: Method for Producing Amide Compounds Using TITLE OF INVENTION: Hydratase from a Thermophilic Bacillus FILE REFERENCE: MSU 4.1-487

CURRENT APPLICATION NUMBER: US/09/549,106

CURRENT FILING DATE: 2000-04-13

PRIOR APPLICATION NUMBER: 60/083,485

PRIOR FILING DATE: 1998-04-29

PRIOR APPLICATION NUMBER: 09/248,528

PRIOR FILING DATE: 1999-02-10

NUMBER OF SEQ ID NOS: 18

SOFTWARE: Patentin Ver. 2.0

SEQ ID NO 2

LENGTH: 1517
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US-09-549-106-2
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APPLICANT:
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NAME/KEY: rRNA
LOCATION: (1).
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                                                                 TYPE: DNA
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APPLICANT: Oriel, Patrick J
APPLICANT: Padmakumar, Rugmini
APPLICANT: Padmakumar, Rugmini
APPLICANT: Kim, Sang H
ITILE OF INVENTION: Method for Producing Amide Compounds Using
ITILE OF INVENTION: Method for Producing Amide Compounds Using
ITILE OF INVENTION: Method for Producing Amide Compounds Using
ITILE REFERENCE: MSU 4.1-488
CURRENT APPLICATION NUMBER: US/09/550,394
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/083,485
PRIOR APPLICATION NUMBER: 60/083,485
PRIOR FILING DATE: 1998-04-29
PRIOR FILING DATE: 1998-04-29
PRIOR FILING DATE: 1999-02-10
NUMBER OF SED ID NOS: 18
SOFTWARE: Patentin Ver. 2.0
SED ID NO 2
LENGTH: 1517
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GENERAL INFORMATION:
APPLICANT: Burrell, Paul
APPLICANT: Blackall, Linda
APPLICANT: Blackall, Linda
APPLICANT: Keller, Jurg
TITLE OF INVENTION: METHOD FOR THE DETECTION OF AQUATIC
TITLE OF INVENTION: NITRITE OXIDISING MICROORGANISMS OF
FILE REFERENCE: CULLN20.001aUS
CURRENT APPLICATION NUMBER: US/09/193,377B
CURRENT FILING DATE: 1998-11-17
NUMBER OF SEQ ID NOS: 62
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 3
LENGTH: 1500
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LOCATION: (1)...(1500)
OTHER INFORMATION: n =
US-09-193-377B-3
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Best Local Similarity 74.8%;
Matches 1121; Conservative
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                                                                                                                                                                                                                           TYPE: DNA
ORGANISM: Nitrospira
FEATURE:
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APPLICANT: Oriel, Patrick J
APPLICANT: Oriel, Patrick J
APPLICANT: Padmakumar, Rugmini
APPLICANT: Aim, Sang H
TITLE OF INVENTION: Method for Producing Amide Compounds Using a l
TITLE OF INVENTION: Hydratase from a Thermophilic Bacillus
FILE REFERENCE: MSU 4.1-401
FILE REFERENCE: MSU 4.1-401
CURRENT APPLICATION NUMBER: US/09/248,528C
CURRENT FILING DATE: 1999-02-10
EARLIER APPLICATION NUMBER: 60/083,485
EARLIER APPLICATION NUMBER: 05083,485
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Best Local Similarity
Matches 1124; Conserv
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ORGANISM: Bacillus
FEATURE:
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CURRENT FILING DATE: 2000-04-13
PRIOR APPLICATION NUMBER: 60/083,485
PRIOR FILING DATE: 1998-04-29
PRIOR APPLICATION NUMBER: 09/248,528
PRIOR FILING DATE: 1999-02-10
NUMBER OF SEQ ID NOS: 18
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Best Local Similarity 75.0
Matches 1124; Conservative
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FEATURE:
NAME/KEY: FRNA
LOCATION: (1)..(1516)
PUBLICATION INFORMATION:
DATABASE ACCESSION NUMBER: Z26930/GenBank
DATABASE ENTRY DATE: 1997-05-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Oriel, Patrick J
APPLICANT: Kim, Sang H
TITLE OF INVENTION: Method for
TITLE OF INVENTION: Hydratase f
FILE REFERENCE: MSU 4.1-486
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AQ960400 LERPD51TR
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	FEATURES Source									COMMENT	JOURNAL		TITLE		AUTHORS	REFERENCE			ORGANISM	SOURCE	KEYWORDS	VERSION	ACCESSION	DEFINITION	Locus	RESULT 1 AQ960400/c	
<pre>/organism="Arabidopsis thaliana" /strain="Landsberg erecta" /db_xref="taxon:3702"</pre>	Location/Qualifiers 1800	Class: shotgun.	Seq primer: TR	Similar to A. thaliana chloroplast sequence (GB:AP000423)	Email: at@tigr.org	Fax: 301 838 0208	Tel: 301 838 0200	9712 Medical Center Dr., Rockville, MD 20850, USA	The Institute for Genomic Research	Contact: Xiaoving Lin	Unpublished (2000)	bolymorphisms	Genomic survey sequencing of Landsberg erecta ecotype of	Feldblyum, T., Liang, F., Creasy, T. and Fraser, C.M.	Buell, C.R., Lin, X., Pai, G., Barnstead, M., Bowman, C., Utterbach, T.,	1 (bases 1 to 800)	Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.	spermatophyta: Wagnoliophyta: endicotyledops: core endicots:	Arabidopsis thaliana	thale cress.	GSS.	AQ960400.1 GI:6788101	sequence. AQ960400	LERFD51TR LERA Arabidopsis thaliana genomic clone LERFD51, DNA	AQ960400 800 bp DNA GSS 28-JAN-2000		

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844 ctatgctcgcaagagtgaaactcaaaggaattgacgggggtccgcacaagcggtggagca
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                                                        CGCTGTGCGTATCGACCCGTGCAGTGCTGTAGCTAACGCGTTAAGTATCCCGCCTGGGGA
                                                                           tgttgggggttttaaccc--tcagtaacgaacctaacggattaagtagaccgcctgggga
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tomato demethylated genomic DNA
Insert Length: 1270 Std Error: 0.00
Seg primer: M13F-R
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Clemson University
100 Jordan Hall, Clemson, SC
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/dev_stage="12-14 weeks post harvest"
/lab_host="E.coli_JM109"
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/db_xref="taxon:4081"
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/clone_lib="cTOG"
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Unpublished (2000)
Contact: Xiaoying Lin
The Institute for Genomic R
9712 Medical Center Dr., Ro
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For addtional_information,
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/clone="LERAP36"
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Best Local Similarity

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Spermatophyta; Magnollophyta; Liliopsida; Pc;
Triticeae; Hordeum.
1 (bases 1 to 868)
Wing,R.; Close,T.J., Kleinhofs,A., Wise,R.,
Y. Anderson,H., Dale,J., Henry,D., Kernodl
                                                                                                      EST.
barley.
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HVSMEC0020H16f Hordeum vulgare seedlin
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HVSMEC0020H16f, mRNA sequence.
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Eukaryota; Viri
  Development of
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gaaggogggatgacgtcaaatcctcatggcctttatgtccaggggccacacacgtgctac
                                                   CGTGTTTAGTTGCCACTATGAGTTTGGAACCCTGAACAGACCGCCGGTGTTAAGCCGGAG
                                                                     tatcgtatgttgctaccttaagttgggcactggtacgaaactgccggtgacaaaccggag
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seq primer: AATTAACCCTACTAAAGGG
High quality sequence start: 16
High quality sequence stop: 700.
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Clemson University
Clemson University
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Unpublished (2000)
On Dec 18, 2000 this sequence version replaced gi:11884381.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /db_xref="taxon:4513"
/clone="HVSMEC0020H16f"
/clone_lib="Hordeum vulgare seedling shoot EST library
HVcDNA0003 (Etiolated and unstressed)"
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/lab_host="TJC121"
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Pred. No. 1.3e-102;
0; Mismatches 178;
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Вb Qy

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27.8%; nilarity 76.9%; Conservative

Score 411; DB 13; Pred. No. 1.6e-102; 0; Mismatches 160;

Length 730;

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599 tacaagtctggagtttgggagaggcaagtggaattccaggtgtagcggtgaaatgcgtag 658 730 TACCCAGCTGGAGTACGGTAGGGGCAGAGGGAATTTCCGGTGGAGCGGTGAAATGCGTAG

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TDGEQ40TH
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Clemson University
100 Jordan Hall, Clemson, SC 29634, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Tomato Demethylated Genomic Unpublished (2001)
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tomato demethylated genomic DNA
Insert Length: 1270 Std Error: 0.00
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                                                                                                                                                                                                                                                                                                                                                                                                                   shotgun
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                                                                                                                                                                                                                                                                         /clone_lib="cTOG"
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               Direct Submission
Direct Submission
Submitted (25-JUN-1999) Genoscope -
BP 191 91006 EVRY cedex - FRANCE (E-
- Web : www.genoscope.cns.fr)
Location/Qualifiers
                                                                                                                                                                                                                                            CNS00WCV 720 bp DNA Arabidopsis thaliana genome of TAMU library from strain genomic survey sequence.
                                                                                             Unpublished 2 (bases 1
                                                                                                                1 (bases 1 to 720)
Salanoubat, M., Choisne, N., Artiguenave, F., Brottier, P., Samson, D., Saurin, W., Weissenbach, J. and Quetier, F.
                                                                                                                                                 Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Sociales; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
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Eukaryota; Viridiplantae;
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∕organism="Arabidopsis
                                                                                                                                                                                                                       GI:5294575
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 thaliana
                                              (E-mail :
                                                         Centre
                                              National de Sequencage 
segref@genoscope.cns.fr
                                             segref@genoscope.cns
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                                                                                                                            ggtcgccaactcgcaagagggagctaatctctaaaagtcggtcccagttcggattggggt 1262
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BH016730
BH016730.1
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TDGCP25TH
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Lycopersicon
Eukaryota; Vi
                    tomato
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/clone="TilH14"
/note="end: SP6"
a 175 c 230 g
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Viridiplantae;
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Pred. No. 1.6e-101;
D; Mismatches 156;
 Streptophyta;
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                                                                                                                                                                                                                      TAGGCGCTGTGCGTATCGACCCGTGCAGTGCTGTAGCTAACGCGTTAAGTATCCCGCCTG
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cagetegtgtegtgagatgttgggttaagteeegeaacgagegeaacceetategtatgt 1076
                                                                               atcatgt-agagatatatgagccttcgggcagatt--cacaggtgctgcatggttgtcgt
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                                                           TCCTCTTGAAAGAGAGGGGTGCCTTCGGGAACGCGGACACAGGTGGTGCATGGCTGTCGT
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Email: http://www.genome.clemson.edu/orders/index.html
tomato demethylated genomic DNA
Seq primer: M13F-R
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/dev_stage="12-14 weeks post harvest"
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/lab_host="E.coli JM109"
/lab_host="E.coli JM109"
/lab_host="Nector: pBluescript SK(-); Site_1: EcoRI digested Stage of the genome of Lycopersicon esculentum ligated fragments of the genome of Lycopersicon esculentum ligated into pBS (SK-). The fragments were cloned into the methylation restrictive E.coli strain JM109 with the purpose of enriching the library for non-methylated DNA fragments. This procedure may enrich the pool of cloned fragments in JM109 cells for sequences representing expressed genes. Average insert size 1.27 kb."

38 a 236;c 190 g 191 t
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/clone="cTOG17F2"
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/cultivar="E6203"
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Pred. No. 1.7e-97;
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cggatagtcctgttggatcacaagatttgataggtaaagatttattgcttggagatgagc
                                      AGTAATATATCGGAACGTGCCTTGTAATGGGGGGATAACTAGTCGAAAGATTAGCTAATAC
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AQ957362
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: Xiaoying Lin
The Institute for Genomic Research
9712 Medical Center Dr., Rockville,
Tel: 301 838 0200
Fax: 301 838 0208
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Buell, C.R., Lin, X., Pai, G., Barnstead, M., Bowman, C., Utter
Feldblyum, T., Liang, F., Creasy, T. and Fraser, C.M.
Genomic survey sequencing of Landsberg erecta ecotype of
Arabidopsis thaliana and identification of sequence-based
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AQ957362 746 bp DNA GSS 28-JAN-2000 LERAP36TF LERA Arabidopsis thaliana genomic clone LERAP36, DNA
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                                                                                                                                                                                                                                                                                                                                                                                                            Email: at@tigr.org
For addtional information,
Seq primer: TF
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sheared to 0.9-1 Kbp before
a 153 c 225 g 159 t
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/strain="Landsberg erecta"
/db_xref="taxon:3702"
/clone="LERAP36"
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73.0%;
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Pred. No. 1.9e-96;
0; Mismatches 181;
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The Institute for Genomic R
9712 Medical Center Dr., Ro
Tel: 301 838 0200
Fax: 301 838 0208
Email: at@tigr.org
                                                                                                                 Eukaryota; Viridiplantae; Streptophyta: Embryophyta; Tracheophyta; Spermatophyta; Magnollophyta; eudicotyledons; core eudicots; Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.

1 (bases 1 to 741)
Buell,C.R., Lin,X., Pai,G., Barnstead,M., Bowman,C., Utterbach,T., Feldblyum,T., Lin,X., Pai,G., Barnstead,M., Bowman,C., Utterbach,T., Genomic survey sequencing of Landsberg erecta ecotype of Arabidopsis thaliana and identification of sequence-based
                                                                                             Unpublished (2000)
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addtional information, see http://www.tigr.org/tdb/at/at.htmliar to A. thaliana chloroplast sequence (GB:AP000423)
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                                             agtttgggagaggcaagtggaattccaggtgtagcggtgaaatgcgtagatatctggagg
                                                                                                                                                                                                                                                                                              acgtatggtgcaagcgttgttcggaatcattgggcgtaaagggtgcgtaggcggatttgt
                                                                                                                                                                                                                                                                                                                                                                                                     gtagtaaacgggattagataccccggtaatccacgccctaaacgttgtctaccagttgtt
                                                                                                                                                                                                                                              AAGTCCGCCGTCAAATCCCAGGGCTCAACCCTGGACAGGCGGTGGAAACTACCAAGCTTG
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Class: shotgun
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/clone_lib="LERA"
/note="Organ: Leaf; Vector: pHOS1; Total genomic sheared to 0.9-1 Kbp before ligation."
a 240 c 169 g 195 t
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/strain="Landsberg erecta"
/db_xref="taxon:3702"
/clone="LERAO33"
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DEFINITION

AQ957182 LERAO34TR

LERA

757 bp DN Arabidopsis

thaliana

genomic

clone

LERAO34, 28-JAN-2000

sequence

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Unpublished (2000)
Contact: Xiaoying Lin
The Institute for Genomic Research
9712 Medical Center Dr., Rockville, 1
Tel: 301 838 0200
Fax: 301 838 0208
                                                                                                                                                                                                                                                                                                                           gcttggagatgagcccgcgcgattagctagttggtgaggtaatggctcaccaaggcga 249
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AQ957182.1
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                         agtttgggagaggcaagtggagttccaggtgtagcggtgaaatgcgtagatatctggagg 669
                                                                                                         AAGTCCGCCGTCAAATCCCCAGGGCTCAACCCTGGACAGGCGGTGGAAACTACCAAGCTTG
AGTACGGTAGGGCAGAGGGAATTTÇCGGTGGAGCGGTGAAATGCGTAGAGATCGGAAAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Trachec Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis 1 (bases 1 to 757)
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For addtional information, see http://www.tigr.org/tdb/at/at.html
Similar.to A. thaliana chloroplast sequence (GB:AP000423)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Seq primer: TR
Class: shotgun.
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/note="Organ: Leaf; Vector: pHOS1; Total genomic force of the color of the col
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/strain="Landsberg erecta"
/db_xref="taxon:3702"
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Pred. No. 1.
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Best Local Similarity
Matches 515; Conserv
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aacaccagtggcgaaggcgacttgctggctcaaaactgacgctgaggcacgaaagcgtgg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Arabidopsis thaliana
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnollophyta; eudicotyledons; core eudicots;
Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
1 (bases 1 to 676)
Rounsley,S.D., Field,C.E., Bass,S., Linher,K., Linher,K., Golden,K., Berry,K., Granger,D., Suh,E., Wible,C., Adams,M.D. and Venter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GSS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            T27K4TR TAMU
                                                                                                                                                                                                                                                                                                                                                                            Department of Eukaryotic Genomics The Institute for Genomic Research 9712 Medical Center Dr., Rockvillu Tel: 301 838 0200
Fax: 301 838 0208
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A BAC End Sequence Database for Identifying Minimal Overlaps in Arabidopsis Genomic Sequencing. Update 3 Uppublished (1997)
                                                                                                                                                                                                                                                                                                                                    Email: rounsley@tigr.org
Seq primer: M13 Reverse
Class: BAC ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Other_GSSs: T27K4TF
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                                                                                                                                                                                                                                                                                                            High quality sequence stop: |
Location/Qualifiers
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                                                                                                                                                                             170
                                                                                            Conservative
                                                                                                                                                                             a
                                                                                                                                                                                                                                                /organism="Arabidopsis
/strain="Columbia"
/db_xref="taxon:3702"
/clone="T27K4"
                                                                                                                                                                         /sex="hermaphrodite"
/note="Vector: BeloBACII; Site_1: HindIII;
; Produced by Rod Wing"
164 c 213 g 128 t 1 others
                                                                                                                                                                                                                               /clone_lib="TAMU"
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Pred. No. 4.8e-93;
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Rockville,
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T27K4, DNA
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RESULT 12
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                                                                                                       Contact: CUGI
Clemson University Genomics
Clemson University
                                       tomato demethylated genomic Seq primer: M13F-R Class: shotgun
                                                                                                                                                                                                                        Lycopersicon esculentum
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
Asteridae; euasterids I; Solanales; Solanaceae; Solanum;
                                                                                                                                                                                                                                                                                                                            sequence.
BH016557
                                                                                                                                                                                                                                                                                                                                                     TDGCN35TH
                                                                                                                                                                                                                                                                                                                                                                   вн016557
                                                                               100 Jordan Hall, Clemson, SC 29634, USA
Email: http://www.genome.clemson.edu/orders/index.html
                                                                                                                                              Unpublished
                                                                                                                                                            Tomato Demethylated
                                                                                                                                                                                       van der Hoeven, R.,
                                                                                                                                                                                                                                                                                                               вн016557.1
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                                                                                                                                                                                                    (bases 1 to 655)
                         Location/Qualifiers
/organism="Lycopersicon
                                                                                                                                                                                                                                                                                                                                                       cTOG
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                                                                                                                                                                                                                                                                                                                                                    655 bp DNA
Lycopersicon
                                                                                                                                                                                      Sun, H.,
                                                                                                                                                            Genomic
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ccagtggcgaaggcgacttgctggctcaaaactgacgctgaggcacgaaagcgtgggtag
                                                                    aattcgatgataccccaaaaacctcacctgggcttgacatggatctgaatc-atgtagag
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                                                       CGATCCCGCGAGGGTGAGCTAACCCCAAAAACCCGTCCTCAGTTCGGATTGCAGGCTGCA
                                                                                                                                                                  GTTTGGAACCCTGAACAGACTGCCGGTGATAAGCCGGAGGAAGGTGAGGATGACGTCAAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /clone="cTOG17E22"
/clone_lib="cTOG"
/tissue_type="young leaves"
/dev_stage="12-14 weeks post harvest"
/lab_host="E_coli_JM109"
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77.08;
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Pred. No. 8.4e-90;
0; Mismatches 142;
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AQ956298/c LOCUS DEFINITION

AQ956298 693 bp DNA LERAJ19TR LERA Arabidopsis thaliana

genomic clone

LERAJ19,

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Hest Local Similarity 77.2%;
Matches 477; Conservativo
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                                                                                                                     CAGTAGTCCTAGCCGTAAACGATGGATACTAGGCGCTGTGCGTATCGACCCCGTGCAGTGC
                                                                                                                                                                                                     gctggctcaaaactgacgctgaggcacgaaagcgtgggtagtaaacggggattagataccc 752
                                                                                                                                                                                                                                                   tccaggtgtagcggtgaaatgcgtagatatctggaggaacaccagtggcgaaggcgactt 692
                                                                                                                                                                                                                                                                                                                         CTCAACCCTGGACAGGCGGTGGAAACTACCAAGCTTGAGTACGGTAGGGGCAGAGGGAAT
                                                                                                                                                                                                                                                                                                                                         ctcaacccgtggcctgcacttgaaactacaagtctggagtttgggagaggcaagtggaat 632
                                                                                                                                   cggtaatccacgccctaaacgttgtctaccagttgttgggggtttttaaccc--tcagtaa 810
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gaattgacgggggtccgcacaagcggtggagcatgtggtttaattcgatgataccccaaa 930
                                                                   cgaacctaacggattaagtagaccgcctggggactatgctcgcaagagtgaaactcaaag 870
                                                                                                                                                                                       GCTGGGCCGACACTGACACTGAGAGACGAAAGCTAGGGGAGCGAATGGGATTAGATACCC
                                                     TGTAGCTAACGCGTTAAGTATCCCGCCTGGGGAGTACGTTCGCAAGAATGAAACTCAAAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Arabidopsis thaliana Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Buell, C.R., Lin, X., Pai, G., Barnstead, M., Bowman, C., Utterbach, T., Feldblyum, T., Liang, F., Creasy, T. and Fraser, C.M. Genomic survey sequencing of Landsberg erecta ecotype of Arabidopsis thaliana and identification of sequence-based
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             9712 Medical Center Dr.,
Tel: 301 838 0200
Fax: 301 838 0208
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Email: at@tigr.org
For addtional information, see http://www.tigr.org/tdb/at/at.html
Similar to A. thaliana chloroplast sequence (GB:AF000423)
Seq primer: TR
Class: shotgun
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Unpublished (2000)
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a 221 c 157 g 178 t 1 others
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/db_xref="taxon:3702"
/clone="LERAJ19"
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tgctggctcaaaactgacgctgaggcacgaaagcgtgggtagtaaacgggattagatacc 751 TGCTGGCCCGACACTGACACTGAGAGACGAAAGCTAGGGGAGCGAATGGGATTAGATACC 578

Query Match Best Local

11 Similarity 486; Conser

24.3%;

Score 359.6; DB Pred. No. 2.3e-88 Mismatches

DB 13;

Indels

6;

Gaps

0;

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AUTHORS
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100 Jordan Hall, Clemson, SC 29634, USA
100 Jordan Hall, Clemson, SC 29634, USA
100 Email: http://www.genome.clemson.edu/orders/index.html
100 tomato demethylated genomic DNA
100 Insert Length: 1270 Std Error: 0.00
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Class: shotgun.
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                                                                /note="Vector: pBluescript SK(-); Site_1: EcoRI; Site_2: Nhol; This library was made from short EcoRI digested tragments of the genome of Lycopersicon esculentum ligated into pBS (SK-). The fragments were cloned into the methylation restrictive E.coli strain JM109 with the purpose of enriching the library for non-methylated DNA fragments. This procedure may enrich the pool of cloned fragments in JM109 cells for sequences representing expressed genes. Average insert size 1.27 kb."
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/db_xref="taxon:4081"
/clone="cTOG30K10"
                                                                                                                                                                                                                                                               /tissue_type="young leaves"
/dev_stage="12-14 weeks post harvest"
/lab_host="E.coli_JM109"
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                                                                          7712 Medical Center Dr., Rockville, Tel: 301 838 0200 Fax: 301 838 0208
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Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.

1 (bases 1 to 676)
Buell, C.R., Lin, X., Pai, G., Barnstead, M., Bowman, C., Utterbach, T.,
Feldblyum, T., Liang, F., Creasy, T. and Fraser, C.M.
Genomic survey sequencing of Landsberg erecta ecotype of
Arabidopsis thaliana and identification of sequence-based
           Seq primer: TF
Class: shotgun
                       Email: at@tigr.org
For addtional information, see http://www.tlgr.org/tdb/at/at.html
Similar to A. thaliana chloroplast sequence (GB:AP000423)
Seq primer: TF
                                                                                                                        Contact: Xiaoying Lin
The Institute for Genomic
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/strain="Landsberg erecta"
/db_xref="rtaxon:3702"
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